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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   | length:
| length:
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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US-09-187-330-36
US-09-187-330-22
US-08-324-297A-1
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US-08-324-297A-1
US-09-187-330-23
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US-09-187-330-23
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US-09-187-330-25
US-08-324-297A-6
US-08-324-297A-7
US-08-324-297A-7
US-08-324-297A-7
US-08-324-297A-7
US-09-248-796A-19008
US-08-324-297A-14
US-08-187-330-24
US-08-187-330-24
US-08-187-330-24
US-08-80-300-3
US-08-80-300-1
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US-08-324-297A-2
                                                     US-08-324-297A-2
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                     Query Match
     Best Local Similarity
                                                                                                                                                                                                     CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/871,
FILING DATE: 22-APR-1992
PRIOR APPLICATION NUMBER: US 07/688,
APPLICATION NUMBER: US 07/688,
FILING DATE: 22-APR-1991
ATTORNEY/AGENT INFORMATION:
NAME: Garrett-Wackowski, Eugen
REGISTRATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 1560
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                       TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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APPLICANT: Brenneman, Dou
APPLICANT: Gozes, Illana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94105-1492
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release CURRENT APPLICATION DATA: APPLICATION NUMBER: US/01 FILING DATE: 17-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
ATREET: One Market Plaza, Steuart Street Tower
                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Neurotrophic Peptides of Activity TITLE OF INVENTION: Dependent Neurotrophic Factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: One Market Pl
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                     STRANDEDNESS:
TOPOLOGY: li
                                                                                                                       TYPE: amino acid
                                                                                                                                       LENGTH:
                                                                                                                                       9 amino acids
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                                                                                       linear
   33.3%;
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PCT-US95-16596-1

PCT-US95-16596-3

PCT-US95-16596-3

PCT-US95-16596-5

PCT-US95-16596-5

PCT-US95-16596-5

PCT-US96-11373-2

US-08-467-822-35

US-08-467-822-35

US-08-466-248-35

US-08-461-722-1

US-08-461-722-1

US-08-468-251-1

US-08-468-251-1
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   Score 40;
Pred. No.
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US-08-981-861-2
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   DB 3; Lo
4.1e+05;
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Scoring table:

Title: Perfect score:

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Matches

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Conservative

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Best Local
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LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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APPLICANT: Brenneman, Douglas E.
APPLICANT: Bassan, Merav
APPLICANT: Zamostiano, Rachel
                                               SEQ ID NO 36
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 015280-291200US
CURRENT APPLICATION UNMER: US/09/187,330
CURRENT FILLING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILLING DATE: 1997-02-07
EARLIER FILLING DATE: 1997-02-06
EARLIER FILLING DATE: 1998-02-06
EARLIER FILLING DATE: 1998-02-06
NUMBER: OF SEQ ID NOS: 63
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                                                                                                                   CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILING DATE: 1997-02-07
EARLIER APPLICATION NUMBER: WO PCT/US98/02485
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.0
                                                                                              SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                          APPLICANT: The Government of the United States of America APPLICANT: as represented by the Secretary of the APPLICANT: Department of Health and Human Services TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III) FILE REFERENCE: 015280-291200US
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Bassan, Merav
APPLICANT: Zamostiano, Rachel
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APPLICANT: Brenneman, Do
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence:ADNF-9 active OTHER INFORMATION: peptide antigen
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ORGANISM: Artificial Sequence
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100.0%; Pred. No.
tive 0; Mismatci
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RESULT

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RESULT 4
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                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/871,973
FILING DATE: 22-APR-1992
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/688,087
FILING DATE: 22-APR-1991
ATTORNEY/AGENT INFORMATION:
                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence:ADNF-9 active OTHER INFORMATION: peptide adsorbed onto bovine serum albumin (BSA) OTHER INFORMATION: as antigen
                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                STRANDEDNESS:
TOPOLOGY: li:
MOLECULE TYPE:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 15
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOPTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
STREET: One Market Plaza, Steuart Street Tower
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: One CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 100 les 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/0 FILING DATE: 17-OCT-1994
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                                                                                                                                                                                                                                                                                 TELEPHONE:
                                                                                 Local
                                                                                                                                                                                                               ENGTH:

    Application US/08324297A
    6174862

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                                 41 SALLRSIPA 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94105-1492
                                                              Similarity 100.0%; Pred. No. 0.014; 9; Conservative 0: Microscherold;
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                                                                                                                                                                                               amino acid
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   SALLRSIPA 10
                                                                                                                                                                                                                  10 amino
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                                                                                                                                                                                                                                                                   (415)
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                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                   (415) 543-9600
                                                                                                                                                peptide
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100.0%; Pred. No. 4.1e+05;
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Dependent Neurotrophic Factor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-324-297A-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 16,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 9; Conserv
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APPLICATION UMBER: US/08/324,297A
FILING DATE: 17-OCT-1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/871,973
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US 07/688,087
FILING DATE: 22-APR-1991
FILING DATE: 22-APR-1991
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EARLIER FILING DATE: 1997-02-07
EARLIER APPLICATION NUMBER: WO PCT/US98/02485
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Zamostiano, Rachel
APPLICANT: The Government of the United States of America
APPLICANT: as represented by the Secretary of the
APPLICANT: Department of Health and Human Services
APPLICANT: Department of Health and Human Services
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CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILING DATE: 1998-11-06
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APPLICANT: Bassan, Merav
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                            ZIP: 94105-1492
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Neurotrophic Peptides of Activity TITLE OF INVENTION: Dependent Neurotrophic Factor NUMBER OF SEQUENCES: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Brenneman, Douglas E. APPLICANT: Gozes, Illana
                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Townsend and Townsend and Crew STREET: One Market Plaza, Steuart Street Tower CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
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100.0%; Pr/
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RESULT 7
US-07-871-973A-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application Patent No. 5767240 GENERAL INFORMATION:
             SOFTWARE: Patentin Release #10, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMMBER: US/07/871,973A
FILING DATB: 22-APR-1992
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/688,087
FILING DATE: 22-APR-1991
ATTORNEY/AGENT INFORMATION:
NAME: Garrett-Wackowski, Eugenia
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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NUMBER OF SEQUENCES: 3
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OTHER INFORMATION:
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OTHER INFORMATION:
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                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: San Francisco
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REGISTRATION NUMBER:
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100.0%; Pr
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40 amino acids wherein each amino
acid is independently selected from
the group consisting of naturally
occurring amino acids and mimetics"
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Center, Eighth Floor
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Matches 9; Conservative
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APPLICANT: Brenneman, Do
APPLICANT: Gozes, Illana
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INFORMATION FOR SEQ ID NO:
                                                                                                  TELEFAX: (415) 543-5043
NFORMATION FOR SEQ ID NO:
                                                                                                                               NAME: Garrett-Wackowski, Eugenia
REGISTRATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 15280-178-1US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/871,973
FILING DATE: 22-APR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/688,087
FILING APPLICATION NUMBER: US 07/688,087
                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                    FILING DATE: 22-APR-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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                 TOPOLOGY:
                               STRANDEDNESS
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/324,297A
FILING DATE: 17-OCT-1994
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                                                                  LENGTH:
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                                                 amino acid
TYPE: peptide
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                                                               14 amino acids
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                     linear
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e Market Plaza, Steuart Street Tower
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-- NO: 2:
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Dependent Neurotrophic Factor
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APPLICANT: Gozes, Illana
APPLICANT: Brenneman, Douglas E.
APPLICANT: Bassan, Merav
APPLICANT: Eassan, Merav
APPLICANT: Zamostiano, Rachel
APPLICANT: The Government of the United States of America
APPLICANT: as represented by the Secretary of the
APPLICANT: as represented by the Secretary of the
APPLICANT: Department of Health and Human Services
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (AD)
FILE REFERENCE: 015280-291200US
CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: WS 60/037,404
EARLIER APPLICATION NUMBER: WO PCT/US98/02485
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
CORTMANDE: DATE: 1998-02-06
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US-09-187-330-23
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CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILING DATE: 1997-02-07
EARLIER APPLICATION NUMBER: WO PCT/US98/02485
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ 1D NOS: 63
SOFTWARE: Patentin Ver. 2.0
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Sequence 21, Apr--
No. 6613740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 23
LENGTH: 14
                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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Best Local Similarity
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APPLICANT: The Government of the United States of America
APPLICANT: as represented by the Secretary of the
APPLICANT: Department of Health and Human Services
APPLICANT: Department of Health and Human Services
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gozes, Ill. APPLICANT: Brenneman,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence:active peptide OTHER INFORMATION: of ADNF I hsp60-related sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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PatentIn Ver. 2.0
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Zamostiano, Rachel
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                                                                                                                                                                                                                                                                                                                                                                                                Application US/09187330
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           33.3%; Score 40; DB 4; 100.0%; Pred. No. 0.02;
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Pred. No.
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US-08-324-297A-6
                                                                                                                       US-08-324-297A-6
                                                                      Query Match
Best Local
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Best Local (
                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/871,973
FILING DATE: 22-APR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/688,087
FILING DATE: 22-APR-1991
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Brenneman, Dov
APPLICANT: Gozes, Illana
                                                                                                                                                                                                  TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: (
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence:ADNF OTHER INFORMATION: I/hsp60-related sequence conjugated through the OTHER INFORMATION: Cys residue to Sephadex for affinity OTHER INFORMATION: chromatography
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/324,297A
FILING DATE: 17-OCT-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
STREET: One Market Plaza, Steuart Street Tower
                                                                                                                                                                                                                                                                       TELEPHONE: (415) 543-9600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                     MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                    Local
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les 9; Conserv
                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                     NAME: Garrett-Wackowski, EuREGISTRATION NUMBER: 37,330
41 SALLRSIPA 49
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                                                    Similarity 9; Conserv
                                                                                                                                                                                      amino acid
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100.0%; Pr
                                                                    33.3%; Score 40; DB 3; 100.0%; Pred. No. 0.029;
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Dependent Neurotrophic Factor
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Pred. No.
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                                                  Mismatches
                                                                                  DB 3; Length 19;
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0.022;
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                                                    Indels
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US-08-324-297A-5
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                                                                                                                                                                                                                       RESULT 13
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; OTHER INFORMATION: /label= Pep277S
US-08-379-613-5
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                                                                                                                                                                Sequence 5, Application US/08324297A Patent No. 6174862
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                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                      GENERAL INFORMATION:
APPLICANT: Brenneman, Douglas E.
APPLICANT: Gozes, Illana
TITLE OF INVENTION: Neurotrophic Peptides of Activity
TITLE OF INVENTION: Dependent Neurotrophic Factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (212)869-8864/9741
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/379,613
FILING DATE: 22-FEB-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Lestie
REGISTRATION NUMBER: 18,72
REGERENCY NOWEER: 18,72
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSES: Townsend and Townsend and Crew
STREET: One Market Plaza, Steuart Street Tower
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 75
TELECOMMUNICATION INFORMATION:
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
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1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24 amino acids
                                                                                                                                                                                                                                                                                                                                                 33.3%; Score 40; larity 100.0%; Pred. No. Conservative 0; Mismatc
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Mismatches
                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 24;
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CITY: San Francisco STATE: California

COUNTRY:

94105-1492

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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/324,297A
FILING DATE: 17-CCT-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: US 07/871,973
APPLICATION NUMBER: US 07/871,973
APPLICATION NUMBER: US 07/871,973
FILING DATE: 22-APR-1992
PRIOR APPLICATION NUMBER: US 07/688,087
APPLICATION NUMBER: US 07/688,087
APPLICATION NUMBER: US 07/688,087
APPLICATION NUMBER: JC-APR-1991
ATTORNEY/AGENT INFORMATION:
NAME: GAITELT-WACKWSki, EUGenia
REGISTRATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 15280-178-1US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
                                                                                                                                               COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: TEM PC COMPATIBLE
COMPUTER: TEM PC COMPATIBLE
COMPUTER: TEM PC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Vers
SOFTWARE: PATENTIAN DATA:
CURRENT APPLICATION DATA:
NDDITCATION NUMBER: US/08/324,297A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (415) 543-5043 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
FILING DALL.

CLASSIFICATION: 514

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/871,973

APPLICATION NUMBER: 22-APR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Brenneman, Douglas E.
APPLICANT: Gozes, Illana
TITLE OF INVENTION: Neuroctrophic Peptides of Activity
TITLE OF INVENTION: Dependent Neurotrophic Factor
NUMBER OF SEQUENCES: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
STREET: One Market Plaza, Steuart Street Tower
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                    APPLICATION NUMBER: US/0 FILING DATE: 17-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH:
                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               42 ALLRSIPA 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30.0%; Score 36; DB 3; L
l Similarity 100.0%; Pred. No. 4.1e+05;
B; Conservative 0; Mismathham
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                                                                                                                                                                                                                                                                                                                                 94105-1492
                                                                                                                                                                                                                                                                                                                                                                                                            San Francisco
                                                                                                                                                                                                                                                                                                                                                                                  California
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                                                                                                                                                                                                              Version #1.30
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FILE REPERBUCE: 107196.132

CURRENT APPLICATION NUMBER: US/09/248,796A

CURRENT FILING DATE: 1999-02-12

PRIOR APPLICATION NUMBER: US 60/074,725

PRIOR FILING DATE: 1998-02-13

PRIOR PILING DATE: 1998-02-13

PRIOR PILING DATE: 1998-08-13

PRIOR FILING DATE: 1998-08-13

NUMBER OF SEQ ID NOS: 28208

SEQ ID NO 19008
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-248-796A-19008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 15
Search completed: February 23, 2005, 19:36:08 Job time : 31.1695 secs
                                                                           片
                                                                                                                                                                                                                                   ; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-19008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICA?
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19008, Apparent No. 6747137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                     Query Match 27.5%;
Best Local Similarity 77.8%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (415) 543-96
TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 22-APR-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                              LENGTH: 409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 15
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                           212 SALLQAIPA
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                                                                                                                41 SALLRSIPA 49
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7; Conserv
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                                                                               220
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Pred. No. 4.1e+05;
2; Mismatches C
                                                                                                                                                                            Score 33; DB 4; Length 409; Pred. No. 32;
                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 9;
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence:
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Searched:
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                   February 23, 2005, 19:34:28; Search time 89 Seconds (without alignments) 327.241 Million cell updates/sec
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120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Published_Applications_AA: *
| Cgn2_6/ptodata/1/pubpaa/USO7_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO9A_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO9B_PUBCOMB.pep:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of score greater than or equal and is derived by analysis is the number of results predicted by chance to have a ater than or equal to the score of the result being printed, rived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query	Length	BU	ID	Description	
,	40	33.3	9	ٔ و	US-09-267-511-1	Sequence 1,	Appli
N	40		9	14	US-10-164-432-3	ω	Appli
w	40		9	15	US-10-296-849-1	Sequence 1,	Appl
4	40		9	15	US-10-623-272-5	Sequence 5,	Appli
U	40		9	15	US-10-623-272-36	36	, Appl
6	40		10	14	US-10-164-432-5	5	Appli
7	40		10	15	US-10-296-849-19		•
80	40		10	15	US-10-623-272-22		•
9	40	33.3	11	15	US-10-296-849-18	Sequence 18	•
10	40		12	15	US-10-296-849-17		, App
11	40		13	15	US-10-296-849-16	Sequence 16	, App
12	40		14	ø	US-09-267-511-21	Sequence 21,	~
13	40		14	_	170		

Conservative

Indels

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Amience 16311	US-10-369-493-162	2 15	w	26.7	32	45
Sequence 15830,	US-10-369-493-158		133	26.7	32	44
Sequence 17371	US-10-437-963-17		787	26.7	32	43
æ	US-10-282-122A-4807		51	•	32	42
e 585	US-10-282-122A-5		47	26.7	32	
11	US-09-815-242-112	9	-	•	32	
m	US-10-437-963-16	9 16	45	٠,	32	39
e 71002,	5 US-10-425-114-71002		394	26.7	32	
e 45954,	US-10-767-701-4		39	26.7	32	37
e 22368,	US-10-369-493-;		32	9	32	
70419,	5 US-10-425-114-70419		288	9	32	35
e 7381,	US-10-032-585-		ود	7.	33	
æ	US-10-	0 15	31	7.	33	33
e 1187	US-10-437-963-11870		163	œ	34	32
e 1187	US-10-437-963-11871		1414	8	34	
e 13	US-10-437-963-134	4 16	1314	28.3	34	30
Sequence 145870	US-10-437-963-14587		88	œ	34	
e 168	US-10-437-963-16896		14	œ	34	
e 139	US-10-437-963-1353		12	8	34	
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e u	US-10-296-849-		8	'n	40	
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e 22,	US-09-267-	9	_	Ψ	40	
e 17,	US-09-267-511-		L	Ψ	40	
e 16,	US-09-267-511-		1-4	'n	40	
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15,	US-09-267-511-		مبو	ü	40	17
equence 21	US-10-623-272-2		_	ω ·	40	
Sequence 23, Ap	US-10-623-272-2		بر	ü	40	15
Ade ter company						

ALIGNMENTS

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GENERAL INFORMATION:

APPLICANT: Brenneman, Douglas B.

APPLICANT: Spong, Catherine Y.

APPLICANT: Spong, Catherine Y.

APPLICANT: Gozes, Illana

APPLICANT: Gozes, Illana

APPLICANT: Bassan, Merav

APPLICANT: The Government of the United States of America

APPLICANT: The Government of the Secretary of the

APPLICANT: as represented by the Secretary of the

APPLICANT: Ramot University Authority for Applied Research

APPLICANT: and Industrial Development, Ltd.

TITLE OF INVENTION: Prevention of Fetal Alcohol Syndrome and Neuronal Cell

TITLE OF INVENTION: Death With ADNF Polypeptides

FILE REFERENCE: 015280-377000US

CURRENT APPLICATION NUMBER: US/09/267,511

CURRENT FILING DATE: 1999-03-12

NUMBER OF SEQ ID NOS: 26
Query Match
Best Local Similarity
The 9; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1
US-09-267-511-1
                                                                                                                                              ; OTHER INFORMATION: Description of Artificial Sequence:activity ; OTHER INFORMATION: dependent neurotrophic factor I (ADNF I) active ; OTHER INFORMATION: site US-09-267-511-1
                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/09267511 Patent No. US20020111301A1
                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                              TYPE: PRT
                           33.3%; Score 40; DB 9; Lo
100.0%; Pred. No. 1.2e+06;
tive 0; Mismatches 0;
                                                                                      Length 9;
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Query Match

DB 15;

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Best Local
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                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/208,944
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 60/267,805
PRIOR PILING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: WO PCT/US01/17758
PRIOR APPLICATION NUMBER: WO PCT/US01/17758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Industrial Development, Ltd.
TITLE OF INVENTION: Use of ADNF for Enhancing Learning and Memory
FILE REFERENCE: 15280W-0042001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Spong, Catherine Y.
APPLICANT: Brenneman, Douglas
APPLICANT: Gozes, Illana
                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                PRIOR FILING DATE: 2001-05-31
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/296,849
CURRENT FILING DATE: 2003-06-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Alcon Inc.
APPLICANT: Clark, Abbot F.
APPLICANT: Debra, Shade L.
                                                       OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Dependent Neurotrophic Factor I (AD) OTHER INFORMATION: core site, SAL or ADNF-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                       TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 09/921,029
PRIOR FILING DATE: 2001-08-02
PRIOR APPLICATION NUMBER: 60/230,964
PRIOR FILING DATE: 2000-09-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 1975A US
CURRENT APPLICATION NUMBER: US/10/164,432
CURRENT FILING DATE: 2002-06-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: The Use of ADNP for the Treatment of Glaucomatous
                                                                                                                                                                                        ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                              ID NO 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            41 SALLRSIPA 49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The Government of the United States of America as represented by the Secretary of the Department of Health and Human Services Ramot University Authority for Applied and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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Pred. No.
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Matches
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)
FILE REFERENCE: 015280-291200US
CURRENT APPLICATION NUMBER: US/10/623,272
CURRENT FILING DATE: 2003-07-17
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/187,330
PRIOR FILING DATE: EARLIER FILING DATE: 1998-11-06
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/037,404
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/037,404
PRIOR FILING DATE: EARLIER FILING DATE: 1997-02-07
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: WO PCT/US98/02485
PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-06
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Publication No. US20040053313A1
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CURRENT FILING DATE: 2003-07-17

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR FILING DATE: EARLIER FILING DATE: 1998-11-06

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR PILING DATE: EARLIER FILING DATE: 1997-02-07

PRIOR PILING DATE: EARLIER FILING DATE: 1997-02-07

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR APPLICATION NUMBER: APPLICATION NUMBER: PRIOR FILING DATE: EARLIER PILING DATE: 1998-02-06

NUMBER OF SEQ ID NOS: 63
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TITLE OF IN
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APPLICANT: Brenn
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9; Conserva
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No. US20040053313A1
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APPLICANT: Brenneman, Douglas
APPLICANT: Gozes, Illana
APPLICANT: The Government of th
APPLICANT: as represented by th
APPLICANT: Department of Health
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SOFTWARE: PatentIn Ver.
SEQ ID NO 36
APPLICANT: The Government of the United States of America APPLICANT: as represented by the Secretary of the APPLICANT: Department of Health and Human Services APPLICANT: Ramot University Authority for Applied and APPLICANT: Industrial Development, Ltd.

TITLE OF INVENTION: Use of ADNF for Enhancing Learning and FILE REFERENCE: 15280W-004200US
CURRENT APPLICATION NUMBER: US/10/296,849
CURRENT FILING DATE: 2003-06-18
PRIOR APPLICATION NUMBER: US 60/208,944
PRIOR FILING DATE: 2000-05-31
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Publication No. US20030166544A1
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Best Local (
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APPLICANT: Debra, Shade L.
TITLE OF INVENTION: The Use of ADNP for the Treatment of Glaucomatous Optic Neuropath
FILE REFERENCE: 1975A US
CURRENT APPLICATION NUMBER: US/10/164,432
CURRENT FILING DATE: 2002-06-06
PRIOR APPLICATION NUMBER: 09/921,029
PRIOR FILING DATE: 2001-08-02
PRIOR FILING DATE: 2001-08-02
PRIOR APPLICATION NUMBER: 60/230,964
PRIOR FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 9
SOPTWARE: PatentIn version 3.1
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ORGANISM: Artificial Sequence
FEATURE:
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TYPE: PRT
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Local Similarity 100.0%; Pred. No.
hes 9; Conservative 0; Mismatch
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                                                                                                                                                                                                                                                                    SEQ ID NO 22
LENGTH: 10
TYPE: PRT
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Matches
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Matches
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                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 015280-291200US
CURRENT APPLICATION NUMBER: US/10/623,272
CURRENT FILING DATE: 2003-07-17
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR PILING DATE: EARLIER FILING DATE: 1998-11-06
PRIOR PILING DATE: EARLIER FILING DATE: 1997-0100 NUMBER: EARLIER APPLICATION NUMBER: EARLIER APPLICATION NUMBER: EARLIER APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR PILING DATE: EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
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APPLICANT: Brenne
APPLICANT: Bassan
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PRIOR FILING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: WO PCT/US01/17758
PRIOR FILING DATE: 2001-05-31
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                  ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:peptide
OTHER INFORMATION: conjugated through the Cys residue to Sephadex
OTHER INFORMATION: affinity chromatography
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                      41 SALLRSIPA 49
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9; Conser
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                                                                          Conservative
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100.0%; Pred. No.
tive 0; Mismatc
                                                                      33.3%; Score 40; DB
100.0%; Pred. No. 0.
Live 0; Mismatches
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                                                                                        DB 15;
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0.17;
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US-10-296-849-18
Sequence 18, Application US/10296849
Publication No. US20040048801A1
GENERAL INFORMATION:

APPLICANT: Spong,

RESULT 9

Brenneman,

Douglas

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APPLICANT: Industrial Development, Ltd.
TITLE OF INVENTION: Use of ADNF for Enhancing Learning and Memory
FILE REFERENCE: 15280W-004200US
CURRENT APPLICATION UNMBER: US/10/296,849
CURRENT FILING DATE: 2003-06-18
PRIOR APPLICATION NUMBER: US 60/208,944
PRIOR PILING DATE: 2000-05-31
PRIOR PILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 60/267,805
PRIOR APPLICATION NUMBER: US 60/267,805
PRIOR APPLICATION NUMBER: WG 7267,805
PRIOR APPLICATION NUMBER: WG 7267,805
PRIOR APPLICATION NUMBER: WG 80/267,805
PRIOR SEQ ID NOS: 23
SOFTWARE: PatentIN Ver. 2.1
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                                                                OTHER INFORMATION: Description OTHER INFORMATION: polypeptide US-10-296-849-17
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Publication No. US20040048801A1
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APPLICANT: Ramot University Authority for Applied and
APPLICANT: Industrial Development, Ltd.
TITLE OF INVENTION: Use of ADNF for Enhancing Learning and Memory
FILE REFERENCE: 15280W-004200US
CURRENT APPLICATION NUMBER: US/10/296,849
CURRENT FILING DATE: 2003-06-18
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                 Query Match
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PRIOR FILING DATE: 2001-05-31
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.1
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PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 60/267,805
PRIOR FILING DATE: 2001-02-08
Best Local Similarity
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TYPE: PRT
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                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Gozes, Illana
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33.3%;
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Pred. No.
Score 40;
Pred. No.
                                                                                                                  of Artificial Sequence: ADNF I
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  DB 15;
0.21;
                      Length 12;
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US-09-267-511-21
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Publication No.
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FILE REFERENCE: 015280-377000US
CURRENT APPLICATION NUMBER: US/09/267,511
CURRENT FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 26
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 21
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PRIOR FILING DATE: 2001-05-31
NUMBER OF SEQ ID NOS: 23
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TITLE OF INVENTION: Use of ADNF for Enhancing Learning and Memory
FILE REFERENCE: 15280W-004220US
CURRENT APPLICATION NUMBER: US/10/296,849
CURRENT FILING DATE: 2003-06-18
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PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 60/267,805
PRIOR FILING DATE: 2001-02-08
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                                                                                                                 APPLICANT: The Government of the United States of America APPLICANT: as represented by the Secretary of the APPLICANT: Department of Health and Human Services APPLICANT: Ramot University Authority for Applied Research APPLICANT: and Industrial Development, Ltd.
TITLE OF INVENTION: Prevention of Fetal Alcohol Syndrome and Neuronal Cell TITLE OF INVENTION: Death With ADNF Polypeptides
                                                                                                                                                                                                                                                                APPLICANT: Spong, Catherine Y. APPLICANT: Gozes, Illana APPLICANT: Bassan, Merav APPLICANT: Zamostiano, Rachel
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TYPE: PRT
ORGANISM: Artificial Sequence
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No. US20040048801A1
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Gozes, Illana
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100.0%; Pred. No.
tive 0; Mismatcl
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APPLICANT: The Government of the United States of America APPLICANT: as represented by the Secretary of the APPLICANT: Department of Health and Human Services APPLICANT: Department of Health and Human Services APPLICANT: Industrial Development, Ltd.

TITLE OF INVENTION: Use of ADNF for Enhancing Learning and Memory FILE REFERENCE: 15280W-00420US
CURRENT APPLICATION NUMBER: US/10/296,849
CURRENT FILING DATE: 2003-06-18
PRIOR APPLICATION NUMBER: US 60/208,944
PRIOR FILING DATE: 2000-05-31
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: US 60/267,805
PRIOR APPLICATION NUMBER: WO PCT/US01/17758
PRIOR APPLICATION NUMBER: WO PCT/US01/17758
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APPLICANT: Brenneman, Douglas
APPLICANT: Gozes, Illana
APPLICANT: The Government of the
APPLICANT: as represented by th
APPLICANT: Department of Health
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Best Local Similarity 100.0%; Pred. No.
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PRIOR FILING DATE: 2001-08-02
PRIOR APPLICATION NUMBER: 60/230,964
PRIOR FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 1975A US
CURRENT APPLICATION NUMBER: US/10/164,432
CURRENT FILING DATE: 2002-06-06
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APPLICANT: Clark, Abbot F.
APPLICANT: Debra, Shade L.
TITLE OF INVENTION: The Use of ADNP for the Treatment
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ORGANISM: Artificial Sequence
FEATURE:
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Search completed: February 23, Job time : 90 secs

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OTHER INFORMATION: Description of Artificial Sequence:active peptide
OTHER INFORMATION: of ADNF I hsp60-related sequence
US-10-623-272-23
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; LENGTH: 14
; TYPE: PRT
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SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 14
LENGTH: 14
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APPLICANT: Brennes
APPLICANT: Bassan
APPLICANT: Zamost
APPLICANT: The Go
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Publication No.
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                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: BARLIER APPLICATION NUMBER: PRIOR FILING DATE: EARLIER FILING DATE: 1998-11-06 PRIOR PELICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR FILING DATE: EARLIER FILING DATE: 1997-02-07 PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-06 NUMBER OF SEQ ID NOS: 63
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Pred. No.
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C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. strain PCC 7120
C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change 09-Jul-2004
C;Accession: AE2407
R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, A.; Irigu Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Takazawa, M.; Yamada, M.; Yasuda, M.; Tabata DNA Res. 8, 205-213, 2001
A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium A;Reference number: AB1807; MUID:21595285; PMID:11759840
A;Accession: AE2407
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A; Introns: 100/1;
A; Note: M3E9.140
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R;Bevan, M.; Vandenbol, M.; Jallet, C.; Portetelle, D.; Hoheisel, J.; Mewes, H.W.; Maye submitted to the Protein Sequence Database, March 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hypothetical protein M3E9.140 - Arabidopsis thallana
C;Species: Arabidopsis thallana (mouse-ear cress)
C;Date: 23-Apr.1999 #sequence_revision 23-Apr.1999 #text_change 09-Jul-2004
                                                                                                                         A;Cross-references: UNIPROT:Q8YMW1; GB:BA000019; PIDN:BAB76512.1; PID:g17133950; GSPDB:A;Experimental source: strain PCC 7120
                                                                                                                                                                    A; Molecule type: DNA
A; Residues: 1-642 < KUR>
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A;Residues: 1-286 <BEV>
A;Cross-references: UNIPROT:065591; EMBL:AL022223
A;Experimental source: cultivar Columbia; BAC clo
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A;Accession: T05061
                                                                                    A;Gene:
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77.8%;
                   28.3%;
87.5%;
  Score 34; DB Pred. No. 26; 1; Mismatches
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Pred. No. 6.7;
2; Mismatches
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26;
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C;Accession: A59266
R;Liang, Y.; Wang, A.; Belyantseva, I.A.; Anderson, D.W.; Probst, F.J.; Barber, an, T.B.; Fridell, R.A.
Genomics 61, 243-258, 1999
A;Title: Characterization of the human and mouse unconventional myosin XV genes A;Reference number: A59266; MUID:20021762; PMID:10552926
A;Accession: A59266
                                 R;Shirai, M.; Hirakawa, H.; Kimoto, M.; Tabuchi, M.; Kishi, F.; Ouchi, K.; Nuccleic Acids Res. 28, 2311-2314, 2000
A;Title: Comparison of whole genome sequences of chlamydia pneumoniae J138
A;Reference number: A86491; MUID:20330349; PMID:10871362
                                                                                                                                                                                               hypothetical protein CPj0220 [imported] - Chlamydophila pneumoniae (strain J138) c;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae Cplate: 02-Mar-2001 #sequence_revision 02-Mar-2
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A; Residues: 1-430 <S
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A; Residues: 1-3530 <LIA>
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;Date: 23-Mar-2001 #sequence_revision 23-Mar-2001 #text_change 09-Jul-2004
;Accession: C86825
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Pred. No. 2.5e+02;
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Pred. No. 28;
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A; Experimental source: C; Genetics:
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Nature 406, 477-483, 2000
A;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae.
A;Reference number: A82035; MUID:20406833; PMID:10952301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid ABC transporter, permease protein VC1861 [imported] - Vibrio cholerae C;Species: Vibrio cholerae C;Species: Vibrio cholerae C;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004
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                                     A;Cross-references: UNIPROT:Q9KQY6; GB:AE004261; GB:AE003852; NID:g9656382; PIDN:AAF950
A;Experimental source: serogroup O1; strain N16961; biotype El Tor
                                                                                                                    A; Residues: 1-225 <HEI>
                                                                                                                                                     A; Molecule type: DNA
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A;Status: preliminary
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, C., Dodson, R., Gwinn, M., Nelson, W., DeBoy, R., Kolonay, J., McClarty, G., Salzber Nucleic Acids Res. 28, 1397-1406, 2000
A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39. A;Reference number: A81500; MUID:20150255; PMID:10684935
                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:Q9Z8W4; GB:AB001608; GB:AB001363; NID:g4376487; PIDN:AAD183
A;Experimental source: strain CWLD29
A;Experimental source: strain CWLD29
R;Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hickey, C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypothetical protein CP0545 [imported] - Chlamydophila pneumoniae (strains CWL029 and C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004 C;Accession: C72104; B81567
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A;Experimental source: strain J138
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A; Residues: 1-181 <STO>
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A; Residues: 1-181 < REA>
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A; Residues: 1-181 < ARN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: Comparative genomes of Clamydia pneumoniae and C. A;Reference number: A72000; MUID:99206606; PMID:10192388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Kalman, S.; Mitchell, W.; Marathe, R.; Lammel, C.; Fan, Nature Genet. 21, 385-389, 1999
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                                                                                                                                                                                                                                                                                                  A;Accession: B81567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Accession: C72104
                                                                                                                                                                                                         ;Cross-references: GB:AE002213; GB:AE002161; NID:g7189446; PIDN:AAF38367.1; PID:g71894
;Experimental source: strain AR39, HL cells
                                                                                  Matches
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52
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                                                                              l Similarity
7; Conserv
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                                                                         26.7%; 5c.
100.0%; Pr
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100.0%;
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Pred.
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Pred. No.
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Gene: VC1861

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A;ACCess.co. DNA
A;Molecule type: DNA
A;Residues: 1-327 <VOS>
A;Cross-references: EMBL:Z75008; NID:g1420278;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: DNA
A;Residues: 1-327 <BEN>
A;Cross-references: EMBL:X94335; NID:gl262139; PID:gl164947
A;Cross-references: EMBL:X94335; NID:gl262139; PID:gl164947
R;Voss, H.; Benes, V.; Rechmann, S.; Teodoru, C.; Schwager,
submitted to the Protein Sequence Database, July 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probable membrane protein YOR100c - yeast (Saccharomyces cerevisiae)
N;Alternate names: hypothetical protein O3193; hypothetical protein YOR3193;
C;Species: Saccharomyces cerevisiae
C;Date: 09-Mar-1996 #sequence revision 12-Apr-1996 #text_change 19-Apr-2002
C;Accession: S61660; S66985
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S61660
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A;Map position: 1
C;Superfamily: histidine
                                          A;Title: Evidence for lateral gene transfer between Archaea and A;Reference number: A72200; MUID:99287316; PMID:10360571 A;Accession: F72331
                                                                                                                                        C;Accession: F72331
R;Nelson, K.E.; Clayton, R.A.; Gill, S.R.; Gwinn, M.L.; Dodson, R.J.; Haft, D.H.; Hi Garrett, M.M.; Stewart, A.M.; Cotton, M.D.; Pratt, M.S.; Phillips, C.A.; Richardson,
                                                                                                                                                                                                    C;Species: Thermotoga maritima
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004
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F;138-222/Domain: ADP,ATP carrier protein repeat homology <ACP2>
F;236-322/Domain: ADP,ATP carrier protein repeat homology <ACP3>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Keywords: transmembrane protein
F;32-127/Domain: ADP,ATP carrier protein repeat homology <ACP1>
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A;Cross-references: SGD:S0005626
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A; Accession: S66985
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A; Accession: S61660
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A;Description: Nucleotide sequence and analysis of a 130 kb fragment of yeast chromosom
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                                                                                                      Nature 399, 323-329, 1999
                                                                                                                                                                                                                                                        conserved hypothetical protein - Thermotoga maritima
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  ;Molecule type: DNA
                    Status: preliminary
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87.5%;
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Pred. No. 35;
0; Mismatches
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Pred. No. 24;
1; Mismatches
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R;Ng, W.V.; Kennedy, S.P.; Mahairas, G.G.; Berquist, B.; Pan, Leithauser, B.; Keller, K.; Cruz, R.; Danson, M.J.; Hough, I Jung, K.H.; Alam, M.; Freitas, T. Proc. Natl. Acad. Sci. U.S.A. 97, 12176-12181, 2000 A;Authors: Hou, S.; Daniels, C.J.; Dennis, P.P.; Omer, A.D.; lA;Title: Genome sequence of Halobacterium species NRC-1. A;Reference number: A84160; MUID:20504483; PMID:11016950 A;Accession: A84387
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C;Genetics:
A;Gene: TM0813
                                                                                                                                                                                                                                                                                                      oligopeptide ABC transporter ATP-binding [imported] - Halobacterium sp. NRC-1 C;Species: Halobacterium sp. NRC-1 C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004 C;Accession: A84387
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Thermotoga maritima
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004
C;Accession: C72424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligopeptide ABC transporter, ATP-binding protein -
                                                                    A;Gene:
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A;Cross-references: UNIPROT:Q9WXR4;
A;Experimental source: strain MSB8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nature 399,
                                                                                                                    A;Molecule type: DNA
A;Residues: 1-440 <STO>
                                                                                                                                                  A; Status: preliminary
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   Matches
                                 Query Match
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Best Local
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 Similarity 7; Conserv
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 Conservative
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                                                                                                    UNIPROT: Q9HMW0;
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2200; MUID:99287316; PMID:10360571
                 26.7%;
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66.7%;
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                   Score 32; Pred. No.
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   Mismatches
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; Maddocks, D.O
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.G.; Jabl
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exodeoxyribonuclease I (EC 3.1.11.1) - Haemophilus influenzae (strain Rd KW20)
C;Species: Haemophilus influenzae
C;Date: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 09-Jul-2004
C;Accession: E64120
C;Accession: E64120
R;Fleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.F.; Kerlavage, Gocayne, J.D.; Scott, J.; Shirley, R.; Liu, L.I.; Glodek, A.; Kelley, J.M.; Weidman, D.M.; Brandon, R.C.; Fine, L.D.; Fritchman, J.L.; Fuhrmann, J.L.; Geoghagen, N.S.M. Science 269, 496-512, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Venter, A;Title: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd. A;Reference number: A64000; MUID:95350630; PMID:7542800
A;Accession: E64120
A;Status: nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chaperonin hsp60, testis - tobacco budworm
C;Species: Heliothis virescens (tobacco budworm)
C;Species: Heliothis virescens (tobacco budworm)
C;Accession: 91-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-200.
C;Accession: S11035
R;Miller; S.G.; Leclerc, R.F.; Erdos, G.W.
J. Mol. Biol. 214, 407-422, 1990
A;Title: Identification and characterization of a testis-specific isoform of the compact of t
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S11035
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A;Residues: 1-473 <TIGR>
A;Cross-references: UNIPROT:P45188;
A;Cross-references: UNIPROT:P45188;
C;Superfamily: exodeoxyribonuclease
C;Keywords: DNA repair; hydrolase
serine O-acetyltransferase homolog yvfD - Bacillus subtilis
C;Species: Bacillus subtilis
C;Accession: E70037
R;Kunst, F:, Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, R;Kunst, F:, Ogasawara, N.; Moszer, I.; Albertini, B.; Capuano, V.; Carter, C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, C.; Bron, S.; Enrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Fe Nature 390, 249-256, 1997
A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; H. Scetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Molecule type: DNA
A;Residues: 1-569 <JMO>
A;Cross-references: UNIPROT:P25420;
A;Note: the sequence from Fig. 8 is
C;Superfamily: chaperonin groEL
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77.8%;
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Ferrari, E.
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A; Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maueel Y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetellt, Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon, A; Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sckowska, A.; Sero, akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama, T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, F., A, Anthors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.
A, Title: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis. A; Reference number: A69580; MUID:98044033; PMID:9384377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: UNIPROT:P71063; GB:Z99121; GB:AL009126; NID:g2635827; PIDN:CAB15429
A;Experimental source: strain 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Status: preliminary; nucleic acid sequence not shown; translation not A;Molecule type: DNA A;Residues: 1-216 KUN>
                                                                                                                                                                                                                                                                                                                                                                                                       R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y.; M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, L. euss, D.; Nierman, W.C.; White, O.; Bisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter, v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hypothetical protein At2937880 [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004
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A,Reference number: A84420; MUID:20083487; PMID:10617197
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A; Residues: 1-247 <STO>
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Q9QUVO
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Q8RXW5
Q65551
Q8W105
Q8W105
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Q8W106
Q8W106
Q8XSW3
Q64UD6
Q7SW3
Q64UD6
Q7XW3
Q74W3
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Submitted (FBB-2002) to the EMBL/GenBank/DDBJ
EMBL; AY080638; AAL85984.1;
InterPro; IPR000555; Mov34_MFN_PAD1.
Pfam; PF01398; Mov34; 1.
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Hypothetical protein At4g26430 (F
Name=At4g26430;
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; rosids;
eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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EMBL; AL022223; CAA18227.1; -.
EMBL; AL161565; CAB79498.1; -.
PIR; T05061; T05061.
InterPro; IPR003639; Mov34-1.
InterPro; IPR003555; Mov34-MFN_PAD1.
Pfam; PF01398; Mov34; 1.
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Q8W205;
01-MAR-2002
01-MAR-2002
SEQUENCE FROM N.A.
MEDLINE=22088475; PubMed=12093376;
MEDLINE=22088475; PubMed=12093376;
Haas B.J., Volfovsky N., Town C.D., Troukhan
Feldmann K.A., Flavell R.B., White O., Salzbb
"Full-length messenger RNA sequences greatly
                                                                           SEQUENCE FROM N.A.

MEDILINE=21607801; PubMed=11742986; DOI=10.1093/emboj/20.24.7096;
Fu H., Reis N., Lee Y., Glickman M.H., Vierstra R.;

"Subunit interaction maps for the regulatory particle of the 26S proteasome and the COP9 signalosome.";

EMBO J. 20:7096-7107(2001).
                                                                                                                                                                                       Name=CSN6B; Synonyms=At4g26430;
Arabidopsis thaliana (Mouse-car cress).
Bukaryota, Viridiplantae, Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core e
Spermatophyta; Magnoliophyta; eudicotyledons; core e
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CSN complex subunit
protein At4g26430).
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SMART; SM00232; JAB_MPN;
Hypothetical protein.
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Bevan M., Vandenbol M., Jallet C., Portetelle D.,
Mewes H.W., Mayer K.F.X., Lemcke K., Schueller C.,
Submitted (MAR-1998) to the EMBL/GenBank/DDBJ data
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MEDLINE=21558789; PubMed=11701877;
Peng Z., Serino G., Deng X.W.;
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Submitted (DEC-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AF395064; AAL58107.1; -.
EMBL; AF09556; AAM65098.1; -.
EMBL; BT002345; AAM65081.1; -.
EMBL; BT002345; AAN86118.1; -.
Pfam; PF01198; MOV34; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COP9 signalosome subunit Name=CSN6B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.

Brover V., Troukhan M., Alexandrov N., Lu Y.-P., Flavel:
Feldmann K.;
Submitted (MAR-2002) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=3702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arabidopsis thaliana (Mouse-ear cress).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAR-2002
01-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hypothetical SEQUENCE 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               annotation.";
Genome Biol. 3:RESEARCH0029-RESEARCH0029(2002)
                                                                                                                                                                                                                                                                                           Pfam;
                                                                                                                                                                                                                                                                                                   "Molecular characterization of subunit
its role in multifaceted developmental
Plant Cell 13:2393-2407(2001).
EMBL: AF434762; AAL49561.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ProDom;
                                                                                                                                                                                                                                                                       ProDom;
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                                                                                                                                                                                                                                                                    PF01398; Mov34; 1.
m; PD363422; Mov34-1; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; PD363422; Mov34-1; 1.
SM00232; JAB MPN; 1.
etical protein.
CE 317 AA; 35404 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      K., Chan M.M., Chang C.H., Dale J.M., Hsuan V.W., Lee J.M., a C.S., Quach H.L., Tang C.C., Toriumi M., Wong C., Wu H.C., Yuan S., Chen H., Cheuk R., Jones T., Kim C.J., Nguyen M., J., Shinn P., Southwick A., Tripp M.G., Wu T., Davis R.W.,
                                                                                                                                                                                                                                    SM00232; JAB MPN; 1.
CE 318 AA; 35493 MW;
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                                                                                                                                                                   Similarity 7; Conser
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                                                                                                    SSLLRSLPA
                                                                                                                                     SALLRSIPA 49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (TrEMBLrel.
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                     PRELIMINARY;
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                                                                                                                                                                   Score 35; DB Pred. No. 52; 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Last sequence update)
Last annotation update)
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Pred. No.
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                     PRT;
                                                                                                                                                                                                                                      6AA06F21C8AEFA29 CRC64;
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                     157
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                                                                                                                                                                                                                                                                                                                                       6 of the COP9 signalosome a processes in Arabidopsis.";
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RESULT
Q9GBG9
ID Q9
AC Q9
DT 01
DT 01
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Best Local S
Matches 7
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GO; GO:0015746; C:mitochondrial electron transport ch:
GO; GO:0005739; C:mitochondrial electron transport ch:
GO; GO:0005739; C:mitochondrial pick it; IEA.
GO; GO:0016491; F:oxidoreductase activity; IEA.
GO; GO:0006118; F:electron transport; IEA.
GO; GO:0006110; P:transport; IEA.
InterPro; IPR005798; Cytb b6 C.
InterPro; IPR005798; Cytb b6 N.
Pfam; PF00032; Cytochrom B C; 1.
Pfam; PF00032; Cytochrom B N; 1.
Pfam; PF00033; Cytochrom B N; 1.
Pfam; PF00033; Cytochrom B N; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUN-2002 (TrEMBLrel. 2
01-JUN-2002 (TrEMBLrel. 2
05-JUL-2004 (TrEMBLrel. 2
Cytochrome b (Fragment).
Q9GBG9
Q9GBG9;
01-MAR-2001
01-MAR-2001
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.

Munasinghe H.N., Murphy
Submitted (MAR-2002) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. Murphy N.P., Augmansinghe D.H.N., Murphy N.P., Augmansinghe D.H.N., murphy N.P., Augmansinghe D.H.N., eque "Utility of mitochondrial DNA seque systematic studies of Australian fi Cherax (Decapoda: Parastacidae)."; Crust. Biol. 23:402-417(2003).
                                                                                                                                                                                                                                                                                                                                       NON_TER
                                                                                                                                                                                                                                                                                                                                                                                           Electron transport; Heme; Mitochondrion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              +
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bumalacostraca; Eucarida; Deo
Parastacoidea; Parastacidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Arthropoda;
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                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                             Transmembrane;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytochrome c1 and the Rieske protein (By similarity).
-I- SIMILARITY: Belongs to the cytochrome b family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mitted (MAY-2002) to the EMBL/GenBank/DDBJ databases.
mitted (MAY-2002) to the EMBL/GenBank/DDBJ databases.
FUNCTION: Component of the ubiquinol-cytochrome c reductase complex (complex III or cytochrome b-c1 complex), which is a respiratory chain that generates an electrochemical potential coupled to ATP synthesis (By similarity).
COPACTOR: Binds two heme groups non-covalently. Heme 1 (or BL 562) is low-potential and absorbs at about 566 nm (By 552) is high-potential and absorbs at about 566 nm (By 552)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   similarity).
SUBUNIT: The main
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157
 (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
                                                                                                                                                                                                                                              Conservative
                                                                           PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                     AA;
                                                                                                                                                                                                                                                                                                                                                                           Transport.
                                                                                                                                                                    152
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17653 MW;
                                                                                                                                                                                                                                                             28.3%;
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21,
27,
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the EMBL/GenBank/DDBJ
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DNA sequences from
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Last annotation update)
Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                             Score 34; DB
Pred. No. 40;
                                                                                                                                                                                                                                                                                                                     8E7A8A8BDB70FE70 CRC64;
                                                                                                                                                                                                                                                Mismatches
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                                                                             379
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                                                                                                                                                                                                                                                                                                                                                                                               Respiratory chain;
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                                                                                                                                                                                                                                                                                Length 157;
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Best Local S
Matches 7
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Cytochrome c1 and the Rieske protein (By similarit:
-|- SIMILARITY: Belongs to the cytochrome b family.

EMBL; AB051239; BAB18191.1; -

EMBL; AB051239; BAB18191.1; -

HSSP; P00157; 1BE3

GG; GO:0016021; C:integral to membrane; IEA.
GG; GO:0005746; C:mitochondrial electron transport cha
GG; GO:0005739; C:mitochondrial; IEA.
GG; GO:0006810; P:lectron transport; IEA.
GG; GO:0006810; P:transport; 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hosoda T., Suzuki H., Harada M., Tsuchiya K., Han S.H., Zhang Y.,
KIYUKOV A.P., Lin L.K.;
"Evolutionary trends of the mitochondrial lineage differentiation in
species of genera Martes and Mustela.";
Genes Genet. Syst. 75.259-267 (2000).
-1- FUNCTION: Component of the ubiquinol-cytochrome c reductase
complex (complex III or cytochrome b-cl complex), which is a
respiratory chain that generates an electrochemical potential
coupled to ATP synthesis (By similarity).
-1- COPACTOR: Binds two heme groups non-covalently. Heme 1 (or BL or
b563) is low-potential and absorbs at about 562 nm, and heme 2 (c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytochrome b.
Name=cytb;
Mustela altaica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q8A9X9;
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SEQUENCE FROM N.A.
STRAIN-VPI-5482 / ATCC 29148;
MEDLINE=22550858; PubMed=12663928;
Xu J., Bjursell M.K., Himrod J., De
Chiang H.C., Hooper L.V., Gordon J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-2003 (TrEMBLrel. 24, Creat
01-JUN-2003 (TrEMBLrel. 24, Last
01-OCT-2003 (TrEMBLrel. 25, Last
                                                                                                                                                                                                                                                                                                                                                           Bacteroides thetaiotaomicron.
Bacteria; Bacteroidetes; Bact
                                                                                                                                                                                                                                                                                                                                                                                                                                               OrderedLocusNames=BT0686;
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                                                                                                                                                                                                                                                                        NCBI_TaxID=818;
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nes 7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               similarity).
SUBUNIT: The main
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H., Harada M., T
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Pred. No.
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         Deng
J.I.;
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annotation update)
                                                       DOI=10.1126/science.1080029;
eng S., Carmichael L.K.,
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Best Local
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PRELIMINARY;
Q73MJ6;
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Q73MJ6 (TrEMBLrel. 27, C
05-JUL-2004 (TrEMBLrel. 27, I
05-JUL-2004 (TrEMBLrel. 27, I
Membrane protein, putative
OrderedLocusNames=TDE1512;
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Q64UD6;
25-OCT-2004
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SEQUENCE
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GO; GO:0005215; F:transporter activity;
GO; GO:0006810; P:transport; IEA.
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EMBL; AE016928; AA075793.1; -.
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Bacteria; Bacteroidetes;
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InterPro; IPR006042; Xan ur permease.
Pfam; PF00860; Xan ur permease; 1.
TIGREAMS; TIGR00801; ncs2; 1.
PROSITE; PS01116; XANTH_URACIL_PERMASE;
[2]
SEQUENCE FROM N.A.
STRAIN=ATCC 35405 / DSM 14222;
PubMed=15064399; DOI=10.1073/pnas.0307639101;
PubMed=15064399; DOI=10.1073/pnas.0307639101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORFNames=BF2146;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kuwahara T., Yamashita A., Hirakawa H., Nakayama H
Kuhara S., Hattori M., Hayashi T., Ohnishi Y.,
"Genomic analysis of Bacteroides fragilis reveals
inversions regulating cell surface adaptation.";
Proc. Natl. Acad. Sci. U.S.A. 0:0-0(2004).
EMBL, AP006841; BAD48893.1;
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25-OCT-2004
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                                                                                                                                                  Treponema denticola.
Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae;
                                                                                                                            NCBI_TaxID=158;
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394 AA;
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(TrEMBLrel. 28,
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Pred. No. 1.1e
2; Mismatches
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Pred. No. 1.1e+02;
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RESULT 13
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TICREPAMS; TIGRO0801; ncs2; 1.
PROSITE; PS01116; XANTH_URACIL_PERMASE;
Complete proteome.
SEQUENCE 432 AA; 45695 MW; 2E732D28i
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        flesh-eater.";

Proc. Natl. Acad. Sci. U.S.A. 99:996-1001(2002).

EMBL; AP003190; BAB81211.1; -.

GO; GO:0016020; C:membrane; IEA.

GO; GO:0005215; F:transporter activity; IEA.

GO; GO:0006810; P:transport; IEA.
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01-MAR-2002 (TrEMBLrel. 20, Last sequ
01-JUN-2003 (TrEMBLrel. 24, Last anno
probable uracil permease.
Name=uraA; OrderedLocusNames=CPE1505;
                         01-MAR-2004
01-MAR-2004
                                                                       Q7S970;
01-MAR-2004
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                                                                                                                        Q7S970
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Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
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AE017251; AAS12029.1;
TDE1512; -.
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Pred. No. 1.2e+02;
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Pred. No. 1.2e+02;
1; Mismatches 0;
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Best Local
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Q9L423;
01-OCT-2000
01-OCT-2000
                                                                                                                         PhosSite; Q9L423; ... PhosSite; Q9L423; ... PhosSite; Q9L423; ... Phosphing; IEA. G0; G0:0004672; F:pprotein kinase activity; G0; G0:0006468; F:pprotein amino acid phosph InterPro; IPR011009; Kinase like. InterPro; IPR000719; Prot_kinase.
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Nature 0:0-0(2003).

-!- CAUTION: The sequence shown here
                              Interro; .....
Pfam; PF00069; Pkinase; 1.
ProDom; PD000001; Prot_kinase; 1.
ProTEIN_KINASE_DOM;
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Pfam; PF02515; COA transf 3; 1.
Hypothetical protein.
SEQUENCE 444 AA; 48688 MW;
                                                                                                                                                                                                                                                                                                                STRAIN=PCC 7120;
Gonzalez L., Phalip V., Zhang C.C.;
Submitted (DEC-1999) to the EMBL/GenBank/DDBJ
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Eukaryota; Fungi; Ascomycota; Pezizomycotina;
Sordariomycetidae; Sordariales; Sordariaceae;
                           Kinase.
                                                                                                                                                                                                                                                                                          EMBL; AJ251821; CAB75357.1;
                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
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Bacteria; Cyanobacteria; Nostocales;
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EMBL; AABX01000217; E
HSSP; P77407; 1PQY.
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  SEQUENCE
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  70785 MW;
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Pred. No. 1.2e+02;
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  81B36952C869E6B7 CRC64;
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phosphorylation; IEA
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hotgun (WGS)
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S., Purcell S., Rehman B.,
B., Butler J., Endrizzi M.,
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3) entry v
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RESULT 15
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Search completed: February 23, 2005, 19:34:16 Job time : 125.65 secs
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X MEDLINE-21595285; pubmed=11759840;
X Maneko T., Nakamura Y., Wolk C.P., Kuritz T., Sasamoto S.,
X Matanabe A., Iriguchi M., Ishikawa A., Kawashima K., Kimura T.,
X Matanabe A., Shimpo S., Sugimoto M., Matsuno A., Muraki A.,
X Kishida Y., Kohara M., Matsumoto M., Takazawa M., Yamada M.,
X Asauda M., Tabata S.;
"Complete genomic sequence of the filamentous nitrogen-fixing
T"Complete genomic sequence of the filamentous nitrogen-fixing
Typonobacterium Anabaena sp. strain PCC 7120.";
DNA Res. 8:205-213(2001).
DNA RES. 8:205-
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Best Local
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Best Local Similarity
Matches 7; Conserv
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Anabhena sp. (strain PCC 7120).
Bacteria; Cyanobacteria; Nostocales; Nostocaceae; Nostoc.
NCBI TaxID=103690;
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01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Serine/threonine kinase.
Serine/threonine kinase.
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519 AMLRSIPA 526
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519 AMLRSIPA 526
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Similarity 87.5%;
7; Conservative
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87.5%; Pred. No. 1.8e+02;
tive 1; Mismatches 0; Indels
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Pred. No. 1.8e+02;
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Total number of hits satisfying chosen parameters:
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121
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed,). B derived λ analysis of the total score distribution

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ADNF III	ADNF III	Activity		Inactive	Human ADN	ADNF III	ADNF III	ADNF III	ADNF III			Seq ID 35	Elongated		Elongated	ADNF III	ADNF III	ADNF III	ACCIVICA

ALIGNMENTS

RESULT 1 AAW64677 04-NOV-1998 AAW64677; AAW64677 standard; protein; Homo sapiens. HIV infection Human ADNF-III peptide fragment. (first entry) Ä

Activity dependent neurotrophic factor III: ADNF-III; ADNP; cell death; activity dependent neuroprotective protein; neurone; excito-coxicity; spinal cord; hippocampus; cerebral cortex; cholinergic; beta-amyloid; N-methyl-D-aspartate; Alzheimer's disease; human immunodeficiency virus;

WO9835042-A2

13-AUG-1998.

06-FEB-1998; 98WO-US002485.

07-FEB-1997; 97US-0037404P

(USSH)

US SEC HEALTH & HUMAN SERVICES

Gozes ŗ, Brenneman DE, Bassan M;

WPI; 1998-447239/38.

Activity dependent neurotrophic factor III polypeptide - useful therapeutically to prevent neuronal cell death associated with e.g. HIV infection, excito-toxicity or Alzheimer's disease.

Example 20; Page 75; 121pp; English.

This sequence represents a peptide used in a method which isolates a novel activity dependent neurotrophic factor III, ANNF-III (also known as activity dependent neuroprotective protein, ANNP). ANNF III polypeptides can be used to prevent neuronal cell death, of e.g. the spinal cord, hippocampal, cerebral cortical or cholinergic neurones associated with e.g. HIV infection, excito-toxicity induced by N-methyl-D-aspartate stimulation or beta-amyloid peptide in Alzheimer's disease. The polypeptides can also be combined with a carrier to alleviate learning impairment produced by cholingeric blockage in Alzheimer's patients. The nucleic acids are useful in polypeptide production and to detect ADNF III

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RESULT 2
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Best Local
novel activity dependent neurotrophic factor III, ADNF-III (also known as activity dependent neurotrophic factor III, ADNF-III (also known as can be used to prevent neuronal cell death, of e.g. the spinal cord, hippocampal, cerebral cortical or cholinergic neurones associated with e.g. HIV infection, excito-toxicity induced by N-methyl-D-aspartate stimulation or beta-amyloid peptide in Alzhaimer'a dicord.
                                                                                                                                                                                                                                                               Activity dependent neurotrophic factor III polypeptide - useful therapeutically to prevent neuronal cell death associated with infection, excito-toxicity or Alzheimer's disease.
                                                                                                                                                                                                                   Claim 24; Page 6; 121pp; English.
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The present sequence is the human Activity Dependent Neurotrophic Factor (ADNF) III-8 or NAP peptide sequence. This peptide was synthesised based on the structural homology to ADNF-9 active peptide and hap60. It is also used as an antigen for immunological detection of cloned ADNF III. ADNF III is also called an Activity Dependent Neuroprotective Protein (ADNP). The human gene was mapped to chromosome 20013.2 and is linked to autosomal dominant nocturnal frontal-lobe epilepsy (ADNFLE) gene. It is expressed in the astrocytes, brain and also in foetal lung and endocrine tissues. This sequence has homology to ADNF I and hap60, heat shock protein and PIF1, a DNA repair protein. The ADNF III polypeptides are useful for the treatment of neurological deficiencies and for prevention

(ADNF III) useful in the treatm preventing neuronal cell death. New nucleic acid encoding an activity (ADNF III) useful in the treatment of

dependent neurotrophic factor neurological deficiencies and

for

2000-376491/32

Example; Page 5; 136pp;

English

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                                                                                                   The present invention relates to the treatment of a condition associated with foetal alcohol syndrome (FAS), involving administering an activity dependent neurotropic factor (ADNF). ADNFs of the present invention may also be used to treat neurological deficiencies and prevent neuronal cell death. The present sequence is an ADNF peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of neuronal cell death associated with gpl20, the envelope protein from HIV, N-methyl-D-Aspartic acid (excito-toxicity); tetrodotoxin (blockage of electrical activity); and beta-amyloid peptide, a substance related to neuronal degeneration in Alzheimer's disease. It is useful for the treatment of Huntington's disease, AIDS dementia complex, epilepsy, neuropathic pain syndromes, Parkinson's disease, amyotrophic lateral sclerosis (ALS), mitochondrial abnormalities, Leber's disease, Wernicke's encephalopathy, Alzheimer's disease, homocysteinuria, hyperprolinemia, encephalopathy, Alzheimer's disease, homocysteinuria, hyperprolinemia,
                                                                               Sequence 8
                                                                                                                                                                                     Claim 9;
                                                                                                                                                                                                              administering
                                                                                                                                                                                                                         Treating condition associated with exposed to alcohol in utero or redu
                                                                                                                                                                                                                                                                                            Brenneman DE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Activity dependent neurotrophic factor; ADNF; FAS; foetal alcohol syndrome; gene therapy; neurologica neuronal cell death.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              encephalopathy, Alzheimer's disease, homocysteinuria, hyperprolinemia sulphite oxide disease, Tourette's syndrome, oxidative stress induced neuronal death, Down's syndrome, developmental retardation and learningairments, drug addiction, tolerance and dependency
                                                                                                                                                                                                                                                                                                                       (UYRA-)
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8; Conserv
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llarity 100.0%;
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                                      33.9%;
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                                                                                                                                                                                                           ociated with fetal alcohol syndrome in a subjutero or reducing neuronal death, involves dependent neurotrophic factors I and/or III.
                                                                                                                                                                                    English.
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Matches 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Treating condition associated with fetal alcohol syndrome in a sexposed to alcohol in utero or reducing neuronal death, involves administering activity dependent neurotrophic factors I and/or I
    Synthetic
                                            Activity dependent neurotrophic factor I; ADNF I; ADNF III; body weight; neuronal cell death; Alzheimer's disease; oxidative stress; VIP; vasoactive intestinal peptide; foetal death; foetal alcohol syndrome.
                                                                                                                                  Activity dependent neurotrophic
                                                                                                                                                                                                                           AAB72322;
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(USSH )
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                                                                                                                                                                                                                                                                     AAB72322 standard;
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3. 1.8e+06;
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                                                                                                                                (ADNF III) peptide
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CC (ADNF I) or ADNF III polypeptides. Sequences AAB72325 - AAB72321 represent CC ADNF II active core site peptides. Sequences AAB72322 - AAB72326 represent CC ADNF III active core site peptides. Sequences AAB72322 - AAB72326 represent CC composition containing either ADNF I, ADNF III and a pharmaceutical CC neuronal cell death, e.g. death of spinal cord neurons, in a patient CC infected with a virus, e.g. human immunodeficiency virus (HIV). The CC neuronal cell death is associated with excito-toxicity induced by N-CC methyl-D-aspartate (NMDE) stimulation, which is induced by beta-amyloid CC peptide in an Alzheimer's disease patient, or induced by cholinergic CC blockade. ADNF I, ADNF III and the pharmaceutical composition are also useful for treating oxidative stress in a patient, for reducing a CC condition, such as decreased body weight, decreased brain weight, compared to the condition of th
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                                                                                                                                                                                           antiparkinsonian; tranquili;
antiinflammatory; antidote;
                                                                                                                                                                                                           ADNF; Activity Dependent Neurotrophic Factor; nootropic; neuroprotective; cerebroprotective; antidiabetic; neuroleptic; anticonvulsant; anti-HIV; antiparkinsonian; tranquilizer; antialcoholic; vulnerary; antibacterial;
31-MAY-2001; 2001WO-US017758.
                                             06-DEC-2001.
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                                                                                                                                    Synthetic.
                                                                                                                                                                           NMDA receptor.
                                                                                                                                                                                                                                                                                                    ADNF III polypeptide active core site peptide
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                                                                                                                                                                                                                                                                                                                                                       26-MAR-2002
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US DEPT HEALTH & HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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SERVICES.
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                                                                                                                                                                                                muscular;
                                                                                                                                                                                                                                                                                                    sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention provides a method of improving learning and/or memory in a subject that involves administering pre- or postnatally an Activity Dependent Neurotrophic Factor (ADNF) to the subject. The ADNF polypeptide is an ADNF I and/or an ADNF III polypeptide comprising the core active site sequences ABB07215 or ABB07216. The method is useful for improving learning and/or memory in a subject; for treating a normal or old subject afflicted with neuropathology, Alzheimer's disease, Down's syndrome, normal mental capacity, mental retardation, for the treatment of central metor systems including degenerative conditions affecting the basal ganglia (see ABB07215 for a detailed description of the various conditions that can be treated by using the ADNF polypeptides). The present sequence represents the active core site sequence of the ADNF III
            WPI; 2003-354501/33
                                                                                                                                                                                                                                       Activity Dependent |
                                    Brenneman DE,
                                                                                                                                                                                                                                                       ADNF I; Activity Dependent Neurotrophic Factor I; neuroprotective; anti-HIV; neuroleptic; antiparkinsonian; nootropic; ADNP; NAP; SAL
                                                            (UYRA-)
                                                                                                  12-SEP-2001; 2001US-0322760P.
10-APR-2002; 2002US-0371961P.
                                                                                                                                                                  20-MAR-2003
                                                                                                                                                                                          WO2003022226-A2
                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                             ADNP I active core peptide
                                                                                                                                                                                                                                                                                                                                                 ABR39742;
                                                                                                                                                                                                                                                                                                                                                                        ABR39742 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8
                                                                                                                                       12-SEP-2002; 2002WO-US029146.
                                                                                                                                                                                                                                                                                                                       23-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Use of an activity dependent neurotropic factor for improving leand/or memory in a subject by pre- or post-natal administration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-2000; 2000US-0208944P.
08-FEB-2001; 2001US-0267805P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 51; 80pp; English.
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                                    Castellon
                                                           HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                           Neuroprotective
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100.0%;
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                                  Spong CY,
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RESULT 9
ADAO7953
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Best Local
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The present invention relates to a method glaucomatous optic neuropathy. The method composition comprising a peptide derived
                                                                                                                                                                         Treating glaucomatous optic neuropathy comprising a peptide derived from or reneuroprotective Factor (ADNP).
                                                                                                                                                                                                                                                                                               WPI; 2003-720933/68
                                                                                                                                                                                                                                                                                                                                                     Clark AF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-SEP-2000;
02-AUG-2001;
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                                                                                                                     Claim 2;
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(SHAD/)
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SHADE
                                                                                                                  Page 2; 13pp; English
                                                                                                                                                                                                                                                                                                                                                        Shade
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2001US-00921029.
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                           for preventing and treating comprises administering a
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The present invention describes a method for preventing or treating an autoimmune disease in a subject. The method comprises administering an activity dependent neurotropic factor (ANNF) polypeptide, where the ADNF polypeptide is a member selected from the group consisting of: (a) an ADNF I polypeptide comprising an active core site with the amino acid sequence of SEQ ID NO:1 (ADQ76120); (b) an ADNF III polypeptide comprising an active core site with the amino acid sequence of SEQ ID NO:2 (ADQ76121); and (c) a mixture of the ADNF II polypeptide of part (a) and the ADNF III polypeptide of part (b). ADNF-encoding nucleic acids, host cells, vectors and antibodies used in the methods are also disclosed

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                                                                                                                                                                                                                                                                                                                                                                                                                              autoimmune disease; multiple sclerosië; myasthenia gravis; Guillain-Barre syndrome; systemic lupus erythematosus; Behcet's syndrome; Sjogren's syndrome; rheumatoid arthritis; Hashimoto's disease; hypothyroiditis; primary biliary cirrhosis; mixed connective tissue disease; chronic active hepatitis; Graves' disease; hypothyroiditis; scleroderma; Graves' disease; chronic idiopathic thrombocytopenic purpura; diabetic neuropathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        neuroprotective factor (ADNP). glaucomatous optic neuropathy. from human ADNP.
                                                                                                                                                               Preventing or treating autoimmune diseases, such as multiple sclerosis, rheumatoid arthritis, hepatitis, Graves' disease, scleroderma and septic shock, using an Activity Dependent Neurotropic Factor (ADNF) polypeptide
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(USSH)
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                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                    septic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vasotropic; muscular; CNS; thyromimetic; antithyroid; antirheumatic
antiarthritic; hepatotropic; virucide; dermatological; haemostatic;
antidiabetic; antibacterial; neurotropic factor agonist;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  activity dependent neurotropic factor; ADNF; ADNF III; active core site; immunosuppressive; nootropic; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-OCT-2004
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1.8e+06;
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nootropic, neuroprotective, antiinflammatory, vasotropic, muscular, CNS, thyromimetic, antithyroid, antirheumatic, antiarthritic, hepatotropic, virucide, dermatological, haemostatic, antidiabetic and antibacterial activities, and can be used as neurotropic factor agonists. The methods and compositions of the present invention are useful for the prevention and/or treatment of autoimmune diseases, such as multiple sclerosis, myasthenia gravis, Guillain-Barre syndrome, systemic lupus erythematosus, Behcet's syndrome, Sjogren's syndrome, rheumatoid arthritis, Hashimoto's disease/hypothyroiditis, primary biliary cirrhosis, mixed connective tissue disease, chronic active hepatitis, Graves' disease/hyperthyroiditis, scleroderma, chronic idiopathic thrombocytopenic purpura, diabetic neuropathy and septic shock. The present sequence represents an ADNF III active core site peptide from the

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ADS73609 standard; peptide; 8

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Matches Query Match Best Local

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Sequence 8

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RESULT 11
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ID ADS7737
XX ADS773
XX ADS773
XX ACT 16-DE
XX ADNF
XX ACT 11-M
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                                                                                                      This sequence represents the active core peptide derived from Activibependent Neurotrophic Factor (ADIF) III. This peptide may be used treating or preventing anxiety or depression in a subject. This sequency optionally be extended at either the N- and/or the C-terminals.

ADNF polypeptide of the invention may be encoded by a nucleic acid to the control of the cont
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                              This specification describes the isolation of novel activity dependent connectorphic factor III, ADNF-III (also known as activity dependent connectorphic factor III, ADNF-III (also known as activity dependent connectorphic factor III, ADNF-III (also known as activity dependent connectorphic factor in the spinal cord, and it is prevent neuronal cell death, of e.g. the spinal cord, consider the spinal cord, and it is carbinal cord, ce.g. HIV infection, excito-toxicity induced by N-methyl-D-aspartate constitution or beta-amyloid peptide in Alzheimer's disease. The collection or beta-amyloid peptide in Alzheimer's disease. The conclect card also be combined with a carrier to alleviate learning conclect acids are useful in polypeptide production and to detect ADNF III conclect acids are useful in polypeptide production and to detect ADNF III conclect acids in biological samples, while the antibodies are useful constructed in the specification but is present in the Sequence ID construction and the specification but is present in the Sequence ID construction in the specification but is present in the Sequence ID construction in the specification but is present in the sequence ID construction in the specification but is present in the sequence ID construction in the specification but is present in the sequence ID construction in the specification but is present in the sequence ID construction in the specification but is present in the sequence ID construction in the specification but is present in the sequence ID construction in the sequ
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The disease is selected from a panic disorder, obsessive-compulsive disorder, post-traumatic stress disorder, social phobia, social anxiety disorder, specific phobias, generalized anxiety disorder, Major depression, dysthymia, and bipolar disorder. The NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin binding site(s) is/are used to identify anxiolytic drugs and NAP-tubulin 
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                   The present sequence is the human Activity Dependent Neurotrophic Factor (ADNF) III generic peptide. It consists of the ADNF III-8 or NAP peptide, flanked by N- and C-terminal generic sites, comprising 1-40 amino acids. ADNF III is also called an Activity Dependent Neuroprotective Protein (ADNP). The human gene was mapped to chromosome 20q13.2 and is linked to autosomal dominant nocturnal fromthellose pilepsy (ADNFE) gene. It is expressed in the astrocytes, brain and also in foetal lung and endocrine tissues. This sequence has homology to ADNF I and hsp60, heat shock
                                                                                                                                                                                                                                                                                                                         New nucleic acid encoding an activity (ADNF III) useful in the treatment of preventing neuronal cell death.
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comprising 1-40 residues, where y= one or
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                                                                                                                                                                                                                                                                                                                                                  dependent neurotrophic factor neurological deficiencies and
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Query Match Best Local Similarity

33.9%; 100.0%;

Score Pred.

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RESULT 14
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The present invention relates to the treatment of a condition associated with foetal alcohol syndrome (FAS), involving administering an activity dependent neurotropic factor (ADNF). ADNFs of the present invention may also be used to treat neurological deficiencies and prevent neuronal cedeath. The present sequence is an ADNF peptide. (Updated on 14-MAY-2003)
                                                                                                                                               Claim 10;
                                                                                                                                                                                            Treating condition associated with fetal alcohol syndrome in a subject exposed to alcohol in utero or reducing neuronal death, involves administering activity dependent neurotrophic factors I and/or III.
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                                                                                                                                                           This invention relates to an activity dependent neurotrophic factor I C (ANNF I) or ANNF III polypeptide. Sequences AAB72315 - AAB72321 represent CC ADNF II active core site peptides. Sequences AAB72322 - AAB72326 represent CC ADNF III active core site peptides. ADNF II, ADNF III and a pharmaceutical CC composition containing either ADNF I or ADNF III and a pharmaceutical CC neurons, cerebral cortical neurons and cholinergic neurons, hippocampal CC neurons, cerebral cortical neurons and cholinergic neurons, in a patient CC infected with a virus, e.g. human immunodeficiency virus (HIV). The CC neuronal cell death is associated with excito-toxicity induced by N-CC methyl-D-aspartate (NMDE) stimulation, which is induced by beta-amyloid CC peptide in an Alzheimer's disease patient, or induced by Cholinergic CC useful for treating oxidative stress in a patient, for reducing a CC condition, such as decreased body weight, decreased brain weight, CC decreased level of vasoactive intestinal peptide (VIP) mRNA, and foetal vor
                                                              Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                  Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 16; Page 57; 88pp; English.
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                                 41 NAPVSIPO 48
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NAPVSIPO 10
                                                                  Conservative
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                                                              33.9%; Score 41; DB 4;
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tive 0; Mismatches
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Title:
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfIles1.pep:*
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RESULT 2 US-09-187-330-33 ; Sequence 33, Application US/09187330 ; Patent No. 6613740 ; PAPPLICANT: INFORMATION: ; APPLICANT: Gozes, Illana ; APPLICANT: Bassan, Merav ; APPLICANT: Bassan, Merav ; APPLICANT: Bassan, Merav ; APPLICANT: The Government of the United States of America	Query Match Best Local Similarity Matches 8; Conser 41 NAPVSIPQ	APPLICANT: Als Government of the officer of the APPLICANT: Department of Health and Human Services APPLICANT: Department of Health and Human Services TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF II FILE REFERENCE: 015280-291200US CURRENT APPLICATION NUMBER: US/09/187,330 CURRENT FILING DATE: 1998-11-06 EARLIER APPLICATION NUMBER: US 60/037,404 EARLIER FILING DATE: 1997-02-07 EARLIER FILING DATE: 1997-02-07 EARLIER FILING DATE: 1998-02-06 NUMBER OF SEQ ID NO 6 ISEQ ID	ALIGNMENTS ESULT 1 S-09-187-330-6 Sequence 6, Application US/09187330 Settent No. 6613740 GENERAL INFORMATION: APPLICANT: Gozes, Illana APPLICANT: Brenneman, Douglas E. APPLICANT: Bassan, Merav APPLICANT: Bassan, Merav APPLICANT: Bassan, Merav APPLICANT: Bassan, Merav	28 31 25.6 317 4 US-09-543-681A-4789 Sequence 4789, 30 31 25.6 392 4 US-09-600-099-6 Sequence 7535, 30 31 25.6 459 6 5194375-6 Sequence 12378 Sequence 1338 Sequence 1339 Sequen
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RESULT 4
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NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 34
LENGTH: 13
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SEQ ID NO 33
LENGTH: 10
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CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILING DATE: 1997-02-07
EARLIER APPLICATION NUMBER: WO PCT/US98/02485
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APPLICANT: The Government of the United States of America
APPLICANT: as represented by the Secretary of the
APPLICANT: Department of Health and Human Services
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)
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APPLICANT: Brenneman, Douglas
APPLICANT: Bassan, Merav
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CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILING DATE: 1997-02-07
EARLIER APPLICATION NUMBER: WO PCT/US98/02485
EARLIER FILING DATE: 1998-02-06
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
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OTHER INFORMATION: polypeptide
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APPLICANT: The Government of the United States of America
APPLICANT: as represented by the Secretary of the
APPLICANT: Department of Health and Human Services
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADN
FILE REFERENCE: 015280-291200US
CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER APPLICATION NUMBER: WS 60/037,404
EARLIER APPLICATION NUMBER: WS 60/037,404
EARLIER FILING DATE: 1998-02-07
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin Ver. 2.0
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; OTHER INFORMATION:
US-09-187-330-12
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SEQ ID NO 12
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LENGTH: 15
TYPB: PRT
ORGANISM: Artificial Sequence
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CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILLING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILLING DATE: 1997-02-07
EARLIER FILLING DATE: 1997-02-07
EARLIER FILLING DATE: 1998-02-06
MUMBER OF SEQ ID NOS: 63
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APPLICANT: Brenneman, Douglas
APPLICANT: Bassan, Merav
APPLICANT: Zamostiano, Rachel
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APPLICANT: as represented by the Secryt of the
APPLICANT: Department of Health and Human Services
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)
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   Similarity
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6613740
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   Conservative
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                                                                                                                                                                   Sequence 57, Application US/09187330
Patent No. 6613740
GENERAL INFORMATION:
APPLICANT: Gozes, Illana
APPLICANT: Brenneman, Douglas E.
APPLICANT: Bassan, Merav
APPLICANT: Zamostiano, Rachel
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Best Local S
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                                                            APPLICANT: The Government of the United States of America APPLICANT: as represented by the Secretary of the APPLICANT: Department of Health and Human Services TITLE OF INVENTION: Activity Dependent Neurotrophic Facto: FILE REFERENCE: 015280-291200US
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EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILING DATE: 1997-02-07
EARLIER APPLICATION NUMBER: WO PCT/US98/02485
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
SOPTWARE: Patentin Ver. 2.0
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APPLICANT: as represented by the Secretary of the
APPLICANT: Department of Health and Human Services
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF
FILE REFERENCE: 015280-291200US
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APPLICANT: Brenneman, Dou
APPLICANT: Bassan, Merav
CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
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TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
ORTHER INFORMATION: Description of Artificial Sequence:activity
OTHER INFORMATION: dependent neurotrophic factor III (ADNF III)
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LOCATION: (49)..(88)
OTHER INFORMATION: Xaa = any amino acid,
OTHER INFORMATION: present or absent
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LOCATION: (1)...(40)
OTHER INFORMATION: Xaa = any amino acid,
OTHER INFORMATION: present or absent
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8; Conserv
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; FEATURE:
; OTHER INFORMATION: human activity dependent neurotrophic factor
; OTHER INFORMATION: (ADNF III)
US-09-187-330-32
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US-09-187-330-31
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LENGTH: 726
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Best Local Similarity
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LENGTH: 781
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Patent No. 6
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            APPLICANT: Gozés, Illana
APPLICANT: Bosen, Douglas E.
APPLICANT: Bassan, Merav
APPLICANT: Zamostiano, Rachel
APPLICANT: Zamostiano, Rachel
APPLICANT: The Government of the United States of America
APPLICANT: as represented by the Secretary of the
APPLICANT: Department of Health and Human Services
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor
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APPLICANT: The Government of the United States of America
APPLICANT: as represented by the Secretary of the
APPLICANT: Department of Health and Human Services
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)
FILE REFERENCE: 015280-291200US
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CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILING DATE: 1997-02-07
EARLIER FILING DATE: 1997-02-06
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin Ver. 2.0
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EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
SOPTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
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REFERENCE: 015280-291200US
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                  Factor III (ADNF III)
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APPLICANT: Zamostiano, Rachel
APPLICANT: The Government of the United States of America
APPLICANT: The Government of the Secretary of the
APPLICANT: Department of Health and Human Services
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)
FILE REFERENCE: 015280-291200US
CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: WS 60/037,404
EARLIER APPLICATION NUMBER: WO PCT/US98/02485
EARLIER APPLICATION NUMBER: WO SECONDAMENT OF SECONDAME: 1998-02-06
NUMBER OF SECONDAME: 1998-02-06
NUMBER: Patentin Ver. 2.0
SECONDAME: Batentin Ver. 2.0
Type: Dem
; Sequence 3, Application US/09187330; Patent No. 6613740; GENERAL INFORMATION:
; APPLICANT: Gozes, Illana
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Best Local S
Matches 8
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APPLICANT: Gozes, Illa
APPLICANT: Brenneman,
APPLICANT: Bassan, Mez
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SEQ ID NO 31
LENGTH: 787
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CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILING DATE: 1997-02-07
EARLIER APPLICATION NUMBER: WO PCT/US98/02485
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
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NAME/KEY: PEPTIDE
LOCATION: (1)..(800)
OTHER INFORMATION: translation of H3'
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ORGANISM: Homo sapiens
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Pred. No.
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PATENT FILING DATE: 1998-02-06

EARLIER APPLICATION UNMBER: 05/07/404

APPLICANT: Brenneman, Douglas E.
APPLICANT: Bassan, Merav

APPLICANT: The Government of the United States of America
APPLICANT: The Government of the United States of America
APPLICANT: Bassan, Merav

APPLICANT: Department of Health and Human Services

ITILE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)

FILE REFERENCE: 015280-291200US

CURRENT APPLICATION NUMBER: US/09/187,330

CURRENT APPLICATION NUMBER: US 60/037,404

EARLIER APPLICATION NUMBER: US 60/037,404

EARLIER APPLICATION NUMBER: WO PCT/US98/02485

EARLIER FILING DATE: 1998-02-06

NUMBER OF SEQ ID NOS: 63

SOFTWARE: Patentin Ver. 2.0

LENGTH: 828

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US-09-187-330-55
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                                                                               Matches
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Best Local :
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CURRENT FILING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILING DATE: 1997-02-07
EARLIER FILING DATE: 1997-02-06
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                       TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III FILE REFERENCE: 015280-291200US
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74 NAPVSIPO 81
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                                     41 NAPVSIPQ 48
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                                                                             Conservative
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                                                                                                33.9%; Score 41;
100.0%; Pred. No.
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RESULT 13 US-09-187-330-59

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Best Local Similarity
Matches 8; Conserve
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SEQ ID NO 59
LENGTH: 874
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LENGTH: 1000
TYPE: PRT
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            PEATURE:

NAME/KEY: MOD RES

NAME/KEY: MOD RES
                                                                                                                 NAME/KEY: MOD RES
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                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)
FILE REFERENCE: 015280-291200US
CURRENT APPLICATION NUMBER: US/09/187,330
CURRENT FILLING DATE: 1998-11-06
EARLIER APPLICATION NUMBER: US 60/037,404
EARLIER FILLING DATE: 1997-02-07
EARLIER FILLING DATE: 1997-02-07
EARLIER FILLING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin Ver. 2.0
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APPLICANT: Zamostiano, Rachel
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APPLICANT: Brenneman, Douglas E.
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APPLICANT: Zamostiano, Rachel
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OTHER INFORMATION: H3' human activity dependent neurotrophic factor
OTHER INFORMATION: III (ADNF III) clone
                                                                                                 LOCATION: (801)
OTHER INFORMATION:
                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
LOCATION: (817)
OTHER INFORMATION: Xaa = unknown
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; LOCATION: (980)
; OTHER INFORMATION: Xaa = unknown
US-09-187-330-1
                                                                                                                                  RESULT 15
US-09-364-609-8
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APPLICANT: Gozes, Illana
APPLICANT: Brenneman, Douglas E
APPLICANT: Zamostiano, Rachel
APPLICANT: Gelber, Edgar
APPLICANT: Pinhasov, Albert
APPLICANT: Bassan, Merav
                                                                                      Sequence 8, Application US/09364609A Patent No. 6649411 GENERAL INFORMATION:
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NAME/KEY: MOD RES
LOCATION: (922)
OTHER INFORMATION:
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LOCATION: (967)
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LOCATION: (959)
OTHER INFORMATION:
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OTHER INFORMATION:
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LOCATION: (870)
OTHER INFORMATION:
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LOCATION: (866)
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OTHER INFORMATION:
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LOCATION: (854)
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OTHER INFORMATION:
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LOCATION: (833)
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LOCATION: (821)
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LOCATION: (964)
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LOCATION: (948)
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APPLICANT: Ramot University Authority for Applied Research & APPLICANT: Industrial Development, Ltd.
TITLE OF INVENTION: Methods of Inhibiting Cancer Cells With ADNF III FILE OF INVENTION: Methods of Inhibiting Cancer Cells With ADNF III TITLE OF INVENTION: Antisense Oligonucleotides FILE REFERENCE: 019856-000100US
CURRENT APPLICATION NUMBER: US/09/364,609A
CURRENT FILING DATE: 1999-07-30
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 8
LENGTH: 1102
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: human activity dependent neurotrophic factor III
OTHER INFORMATION: (ADNF III) cDNA
US-09-364-609-8
Search completed: February 23, 2005, 19:36:17 Job time : 38.8305 secs
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                                                                                                                       354 NAPVSIPO 361
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Maximum DB seq length: 2000000000
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121
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/ Cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US10D_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US10D_PUBCOMB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/ Cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
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_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
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6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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41	4 1	41	41	41	41	41	41	41	41	41	41	Score
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15	15	15	13	13	10	10	10	8	8	8	80	Query Match Length
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US-10-623-272-35	115-10-296-849-22	US-10-623-272-34	US-10-296-849-21	US-09-267-511-24	US-10-623-272-33	US-10-296-849-20	US-09-267-511-23	US-10-623-272-6	US-10-296-849-2	US-10-164-432-4	US-09-267-511-2	ID
Sequence 22, Appl			Sequence 21, Appl	Sequence 24, Appl	Sequence 33, Appl		Sequence 23, Appl	Sequence 6, Appli	Sequence 2, Appli	Sequence 4, Appli	Sequence 2, Appli	Description

ALIGNMENTS

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APPLICANT: Spong, Catherine Y.
APPLICANT: Gozes, Illana
APPLICANT: Zamostiano, Rachel
APPLICANT: Zamostiano, Rachel
APPLICANT: The Government of the United States of America
APPLICANT: Bepartment of Health and Human Services
APPLICANT: Department of Health and Human Services
APPLICANT: and Industrial Development, Ltd.
TITLE OF INVENTION: Prevention of Fetal Alcohol Syndrome and Neuronal Cell
TITLE OF INVENTION: Death With ADNF Polypeptides
FILE REFERENCE: 015280-377000US
CURRENT APPLICATION UNMBER: US/09/267,511
CURRENT FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 26
                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence:activity; OTHER INFORMATION: dependent neurotrophic factor III (ADNF III); OTHER INFORMATION: active site US-09-267-511-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1
US-09-267-511-2
                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
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     Matches
                           Query Match
Best Local 9
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  Similarity 100
8; Conservative
33.9%; Score 41; DB 9;
100.0%; Pred. No. 1.2e+C
tive 0; Mismatches
  2e+06;
                                                    Length 8;
     Indels
     0;
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Gaps

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NAPVSIPO 48

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; ORGANISM: Artificial Sequence; FEATURE; OTHER INFORMATION: Description of Artificial Sequence:Activity; OTHER INFORMATION: Dependent Neurotrophic Factor III (ADNF III); OTHER INFORMATION: active core site, NAP or ADNF III-8 US-10-296-849-2
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US-10-296-849-2
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Best Local S
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
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SEQ ID NO 4
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PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 60/267,805
PRIOR FILING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: WO PCT/US01/17758
PRIOR FILING DATE: 2001-05-31
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ramot University Authority for Applied and APPLICANT: Industrial Development, Ltd.
TITLE OF INVENTION: Use of ADNF for Enhancing Learning FILE REFERENCE: 15280W-004200US
CURRENT APPLICATION NUMBER: US/10/296,849
CURRENT FILING DATE: 2003-06-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Spong,
APPLICANT: Brenne
APPLICANT: Gozes,
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PRIOR FILING DATE: 2001-08-02
PRIOR APPLICATION NUMBER: 60/230,964
PRIOR FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 9
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APPLICANT: Clark
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APPLICANT: Debra, Shade L.
TITLE OF INVENTION: The Use of ADNP for the Treatment of Glaucomatous Optic Neuropath RILE REFERENCE: 1975A US
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CURRENT FILING DATE: 2002-06-06
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ORGANISM: homo sapiens
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8; Conserv
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Pred. No.
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1.2e+06;
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GENERAL INFORMATION:
APPLICANT: Brenneman, Douglas E
APPLICANT: Spong, Catherine Y.
APPLICANT: Gozes, Illana
APPLICANT: Bassan, Merav
APPLICANT: Camostiano, Rachel
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Matches
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LENGTH: 8
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APPLICANT: Gozes, I
APPLICANT: Brenner
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                                                                                           APPLICANT: The Government of the United States of America APPLICANT: as represented by the Secretary of the APPLICANT: Department of Health and Human Services APPLICANT: Ramot University Authority for Applied Research APPLICANT: and Industrial Development, Ltd.
TITLE OF INVENTION: Prevention of Fetal Alcohol Syndrome and Neuronal Cell TITLE OF INVENTION: Death With ADNF Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                      Patent No. US20020111301A1
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CURRENT FILING DATE: 2003-07-17

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR FILING DATE: EARLIER FILING DATE: 1998-11-06

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR FILING DATE: EARLIER FILING DATE: 1997-02-07

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: PRIOR FILING DATE: UNMBER: EARLIER APPLICATION NUMBER: PRIOR APPLICATION NUMBER: PRIOR FILING DATE: EARLIER APPLICATION NUMBER: PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-06

NUMBER OF SEQ ID NOS: 63
FILE REFERENCE: 015280-377000US
CURRENT APPLICATION NUMBER: US/09/267,511
CURRENT FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 26
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TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)
FILE REFERENCE: 015280-291200US
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100.0%; Pred. No. 1.
cive 0; Mismatches
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100.0%; Pred. No.
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Query Match
Best Local Similarity
Watches 8; Conserve
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                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Gozes, I
APPLICANT: Brennem
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PRIOR FILING DATE: 2001-05-31
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 20
                                                                                                                                                                                     Sequence 33, Application US/10623272 Publication No. US20040053313A1
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Best Local (
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LENGTH: 10
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TITLE OF INVENTION: Use of ADNF for Enhancing Learning and FILE REFERENCE: 15280W-004200US CURRENT APPLICATION NUMBER: US/10/296,849
CURRENT FILING DATE: 2003-106-18
                                                                                                    APPLICANT:
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PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 60/267,805
PRIOR FILING DATE: 2001-02-08
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APPLICANT: Brenne
APPLICANT: Gozes,
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ORGANISM: Artificial Sequence
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Brenneman, Douy....
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Bassan, Merav
Bassan, Merav
Zamostiano, Rachel
The Government of the United States of America
The Government by the Secretary of the
The Government of Health and Human Services
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Gozes, Illana
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SEQ ID NO 24
LENGTH: 13
TYPE: PRT
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CURRENT FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 26
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TITLE OF INVENTION: Prevention of Fetal Alcohol Syndrome and Neuronal Cell TITLE OF INVENTION: Death With ADNF Polypeptides
FILE REFERENCE: 015280-377000US
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PRIOR FILING DATE: EARLIER FILING DATE: 1998-11-06
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/037,404
PRIOR FILING DATE: EARLIER FILING DATE: 1997-02-07
PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-06
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CURRENT APPLICATION NUMBER: US/10/623,272
CURRENT FILING DATE: 2003-07-17
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ORGANISM: Artificial Sequence
FEATURE:
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RESULT 9 US-10-296-849-21

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LENGTH: 13
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CURRENT FILING DATE: 2003-07-17

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/187,330

PRIOR EILING DATE: EARLIER FILING DATE: 1998-11-06

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/037,404

PRIOR FILING DATE: EARLIER FILING DATE: 1997-02-07

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: WO PCT/US98/0:
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                                                                                                                                                              SOFTWARE: PatentIn
                                                                                                                                                                                                         PRIOR FILING DATE:
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PRIOR APPLICATION NUMBER: US 60/208,944
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 60/267,805
PRIOR FILING DATE: 2001-02-08
PRIOR APPLICATION NUMBER: WO PCT/US01/17758
PRIOR FILING DATE: 2001-05-31
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TITLE OF INVENTION: Use of ADNF for Enhancing Learning and FILE REFERENCE: 15280W-004200US
CURRENT APPLICATION NUMBER: US/10/296,849
CURRENT FILING DATE: 2003-06-18
                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 63
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OTHER INFORMATION: Description OTHER INFORMATION: polypeptide
                                                                TYPE: PRT ORGANISM: Artificial Sequence
                                               FEATURE:
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Similarity 100.0%;
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No. US20040053313A1
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Gozes, Illana
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                   of Artificial Sequence:ADNF III
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Best Local
                                                                    PRIOR APPLICATION NUMBER: US 60/208,944
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: US 60/267,805
                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                               APPLICANT: Spong, APPLICANT: Brenne
                                                                                                                           CURRENT APPLICATION NUMBER: US/10/296,849
CURRENT FILING DATE: 2003-06-18
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CURRENT FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 26
SOFTWARE: Patentin Ver. 2.1
PRIOR APPLICATION NUMBER: WO PCT/US01/17758
PRIOR FILING DATE: 2001-05-31
NUMBER OF SEQ ID NOS: 23
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TYPE: PRT
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TITLE OF INVENTION: Prevention of Fetal Alcohol Syndrome and Neuronal Cell TITLE OF INVENTION: Death with ADNF Polypeptides
                                                         PRIOR FILING DATE: 2001-02-08
                                                                                                                                                                FILE REFERENCE: 15280W-004200US
                                                                                                                                                                               APPLICANT: Industrial Development, Ltd.
TITLE OF INVENTION: Use of ADNF for Enhancing Learning and Memory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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Similarity 100.0%;
8; Conservative
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Gozes, Illana
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Zamostiano, Rachel
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Gozes, Illana
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100.0%; Pred. No. 0.4
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Pred. No.
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Query Match
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Watches 8; Conserve
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GENERAL INFORMATION:
APPLICANT: Brenneman, Douglas E.
APPLICANT: Spong, Catherine Y.
APPLICANT: Gozes, Illana
APPLICANT: Bassan, Merav
APPLICANT: Bassan, Merav
APPLICANT: Zamostiano, Rachel
APPLICANT: The Government of the United States of APPLICANT: as represented by the Secretary of the
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SEQ ID NO 22
LENGTH: 15
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Publication No.
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APPLICANT: The Government of the United States of America
APPLICANT: as represented by the Secretary of the
APPLICANT: as represented by the Secretary of the
APPLICANT: Department of Health and Human Services
TITLE OF INVENTION: Activity Dependent Neurotrophic Factor III (ADNF III)
FILE REFERENCE: 015280-291200US
CURRENT APPLICATION NUMBER: US/10/623,272
CURRENT FILING DATE: 2003-07-17
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/187,330
PRIOR PILING DATE: EARLIER FILING DATE: 1991-11-06
PRIOR FILING DATE: EARLIER FILING DATE: 1997-02-07
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/037,404
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: WO PCT/US98/02485
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: WO PCT/US98/02485
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No. US20040053313A1
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APPLICANT: Ramot University Authority for Applied Research
APPLICANT: and Industrial Development, Ltd.
TITLE OF INVENTION: Prevention of Fetal Alcohol Syndrome and Neuronal Cell
TITLE OF INVENTION: Death With ADNF Polypeptides
FILE REFERENCE: 015280-377000US
CURRENT APPLICATION NUMBER: US/09/267,511
CURRENT FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 26
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 19
                                                                                                                                                                                                                                                               SEQ ID NO 26
LENGTH: 17
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APPLICANT: Zamostiano, Rachel
APPLICANT: Zamostiano, Rachel
APPLICANT: The Government of the United States of America
APPLICANT: The Government of the Secretary of the
APPLICANT: Ramot University Authority for Applied Research
APPLICANT: and Industrial Development, Ltd.
TITLE OF INVENTION: Prevention of Fetal Alcohol Syndrome and Neuronal Cell
TITLE OF INVENTION: Death With ADNF Polypeptides
FILE REFERENCE: 015280-377000US
CURRENT APPLICATION NUMBER: US/09/267,511
CURRENT FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 26
SOFTWARE: Patentin Ver. 2.1
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TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
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        GenCore version 5.1.6
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Q8E377
Q8K7F1
Q8K7F1
Q8K7F1
Q9EZG3
Q9Y7ZB
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Q6zq47 mus musculu
Q9h2p0 homo sapien
Q6dhz8 homo sapien
Q6c9b5 yarrowia li
Q82yw3 enterococcu
                                Q8bqb7
Q6ffk0
Q841k3
Q9v3b2
Q9v3b2
Q6maf9
Q55112
Q7yqm2
Q7yqm1
                                                                                                                                                                                                                                                                                  Q92xp0
Q9xcf3
Q740v0
O28744
                                                                                                                                                                                                                                                                                                                                                                                                Q8k7f1
Q8duc1
Q9i2g3
Q9i2g3
Q6ka80
Q6ka80
Q8bx17
Q8bx17
Q8szd3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q8e338
Q8p0z3
Q99zk7
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1 pongo pygma
1 homo sapien
     homo sapien
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Result No.

ALIGNMENTS

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clear protein (atypical).	E FROM N.A. , Xu K., Du Y.; te sequence of a rat protein containing a femtomolar- nt neuroprotective peptide."; ed (FEB-2000) to the EMBL/GenBank/DDBJ databases. CCTION: Potential transcription factor. May mediate so roprotective peptide VIP-associated effects involving with and cancer proliferation. CCELLULAR LOCATION: Nuclear (Potential). UCTION: By the neuroprotective peptide VIP. ILARITY: Contains 1 homeobox domain. ILS-PROT entry is copyright. It is produced through a the Swiss Institute of Bioinformatics and the EMB opean Bioinformatics Institute. There are no restri non-profit institutions as long as its content d and this statement is not removed. Usage by and s requires a license agreement (See http://www.isb-si an email to license@isb-sib.ch).	

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RESULT
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Matches 8
                                                               EMBL; AF068198; AAD19843.1; -.
MGD; MGI:138758; AADD.
InterPro; IPR001356; Homeobox.
InterPro; IPR009057; Homeodomain_1:
InterPro; IPR007087; Znf_C2H2.
Pfam; PF00046; Homeobox; 1.
Pfam; PF00096; zf_C2H2; 1.
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DNA_BIND
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                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADNP_MOUSE STANDARD; PRT; 828 AA.

(92103;
(28-FEB-2003 (Rel. 41, Created))
28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Activity-dependent neuroprotector (Activity-dependent neuroprotective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-99155106; PubMed=10037502;
Bassan M., Zamostiano R., Davidson
Perl O., Bassan H., Blat C., Gibne
SMART; SM00389; HOX; 1.
SMART; SM00355; ZnF_C2H2; 4.
SMOSITE; BS50071; HÖMEOBOX 2; 1.
PROSITE; PS00028; ZINC_FINGER_C2H2_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10090;
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Complete sequence of a novel protein containing a activity-dependent neuroprotective peptide.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gozes I.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FISSUE=Brain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.,
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                                                                                                                                                                                                                                                                                                                       concentrations against toxicity associated with tetrodoxin (electrical blockade), the beta-amyloid peptide (the Alzheimer's disease neurotoxin), N-methyl-aspartate (excitotoxicity), and the human immunideficiency virus (HIV) envelope protein. SIMILARITY: Contains 1 homeobox domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FUNCTION: Potential transcription factor. May mediate some of the neuroprotective peptide VIP-associated effects involving normal growth and cancer proliferation.
SUBCELLULAR LOCATION: Nuclear (Potential).
                                                                                                                                                                                                                                                                                                                                                                                                      INDUCTION: By the neuroprotective peptide VIP. MISCELLANBOUS: When isolated from the sequence, the neuroprotective peptide provides neuroprotection at subfemtomolar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBCELLULAR LOCATION: Nuclear (Potential).
TISSUE SPECIFICITY: Expressed in the brain, with a higher expression in cerebrellum and hippocampus. Weakly expressed in lung, kidney and intestine, and expressed at intermediate level
                                                                                                                                                                                                                                                                                                                                                                                                                                                          test18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 823 AA;
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534
91335
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AND SYNTHESIS
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C., Gibney G., Glazner G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 41;
Pred. No.
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Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A4C4BC616052DBDF CRC64;
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thi; Muridae;
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Brenneman
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; Murinae; Mus
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Matches 8
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Matches 8
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C 1-1 SUBCELLULAR LOCATION: Nuclear (By similarity).

R GO; GO:0005634; C:nucleus; IEA.

R GO; GO:0003700; F:transcription factor activity; IEA.

R GO; GO:0003700; F:transcription of transcription, DNA-dependence of transcription of transcription, DNA-dep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Transcription
ZN_FING 16
ZN_FING 20
ZN_FING 23
ZN_FING 33
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Q6ZQ47;
Q5_JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Saga Y., Nagase T., Ohara O., Koga H.; "Prediction of the coding sequences of mouse homologues of KIAA gene: III. the complete nucleotide sequences of 500 mouse KIAA-homologous cDNAs identified by screening of terminal sequences of cDNA clones randomly sampled from size-fractionated libraries."; DNA Res. 10:167-180(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
MCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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8; Conser
                                                                                                                                        Similarity
  NAPVSIPO 342
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          188
229
254
366
405
533
81
                                                                                                  33.9%; 50
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         33.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ohara R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WW.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zinc-finger.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Last sequence update)
                                                                                                              Score 41; DB; Pred. No. 17; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>.</u>
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C2H2-type 2.
C2H2-type 3.
C2H2-type 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C2H2-type 3.
C2H2-type 4
C2H2-type 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 41;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Glu-ri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Neuroprotective peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homeobox
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         _C2H2_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9DFE669C506E8606 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Inamoto S.,
                                                                                                                                                                                                                             7FF389C7FAEDF660 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nuclear protein; Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (atypical).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (atypical)
                                                                                                                                                                     2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Koseki H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                 0
                                                                                                                                                                     Length 1089;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA-dependent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 828;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hiraoka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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RA Deloukas P., Matthews L.H., Ashurst J.L., Burton J., Gilbert J.G.R., Almeida J.P., Babbage A.K., Bagguley C.L., RA Bailey J., Barlow K.F., Bates K.N., Beard L.M., Beare D.M., RA Bailey J., Barlow K.F., Bates K.N., Beard L.M., Beare D.M., RA Bailey J., Bird C.P., Blakey S.E., Bridgeman A.M., Brown A.J., RA Beasley O.P., Bird C.P., Blakey S.E., Bridgeman A.M., Brown A.J., RA Chapman J.C., Clamp M., Clark G., Clark L.N., Clark S.Y., Clee C.M., RA Chapman J.C., Clamp M., Clark G., Clark L.N., Clark S.Y., Clee C.M., RA Chapman J.C., Cobley V.E., Collier R.E., Connor R.E., Corby N.R., RA Clegg S., Cobley V.E., Collier R.E., Connor R.E., Corby N.R., RA Clegg S., Cobley V.E., Collier R.E., Connor R.E., Corby N.R., RA Clegg S., Cobley V.E., Cartfiths M.N.D., Gwilliam R., Hall R.E., RA Craffam D.V., Griffiths C., Griffiths M.N.D., Gwilliam R., Hall R.E., RA Hammond S., Harley J.L., Heath P.D., Ho S., Holden J.L., Howden P.J., RA Huckle E., Hunt A.R., Hunt S.E., Jekosch K., Johnson C.M., Johnson D., RA Kay M.P., Kimberley A.M., King A., Knights A., Laird G.K., Lawlor S., RA Kay M.P., Kimberley A.M., King A., Knights A., Laird G.K., Lawlor S., RA Marsh V.L., Martin S.L., McConnachie L.J., McLay K., McMurray A.A., RA Milne S.A., Mistry D., Moore M.J.F., Mullikin J.C., Nickerson T., RA Milne S.A., Nistry D., Moore M.J.F., Mullikin J.C., Nickerson T., RA Philliames B.J.C.T., Prathalingam S.R., Plumb R.W., Ramasy H., RA Swann R.W., Sycamore N., Taylor R., Teeward C.A., Sulston J.E., RA Swuce C.D., Smith M.L., Soderlund C., Steward C.A., Sulston J.E., RA Wille B.G., Wallie M., Wallie J.M., Thorpe A., RA Wille A., Wallie J.M., Wallie S.A., Beck S., RA Wilning L., Warly P.W., Hubbard T., Durbin R.M., Bentley D.R., Beck S., RA Milne B.J., Wallie S.A., Hacker P., Willeys D.L., Williams S.A., Beck S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE-Fetal brain;
TISSUE-Fetal brain;
PubMed=11013255; DOI=10.1074/jbc.M007416200;
Pamostiano R., Pinhasov A., Gelber E., Steingart R.A.,
Zamostiano R., Wollman Y., Eyre H.J., Mulley J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q9H2P0; O94881; Q9UG34;
28-FEB-2003 (Rel. 41, C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation updat
              Wambutt R., Heubner D., Mewes H.-W., Gassenhuber J., Wiemann S.;
Submitted (JUN-1999) to the EMBI/GenBank/DBJ databases.
-!- FUNCTION: Potential transcription factor. May mediate some of the
neuroprotective peptide VIP-associated effects involving normal
                                                                                                                                                                                                                                                                                                                                                                                 Rogers J.; "The DNA sequence and comparative analysis of human chromosome 20."; Nature 414:865-871(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-2004 (Rel. 44, Last annotation update)
Activity-dependent neuroprotector (Activity-dependent neuroprotective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HUMAN
                                                                                                                                                                         "Prediction of the coding sequences of unidentified human genes. The complete sequences of 100 new cDNA clones from brain which co for large proteins in vitro.";
DNA Res. 5:277-286(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Brenneman D.E., Gozes I.; "Cloning and characterization of the human activity-dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein)
                                                                                                         rissum=Uterus;
                                                                                                                               SEQUENCE OF
                                                                                                                                                                                                                                                                Nagase T., Ishikawa K.-I., Suyama M., Kiku
Tanaka A., Kotani H., Nomura N., Ohara O.;
                                                                                                                                                                                                                                                                                                             MEDLINE=99087487; PubMed=9872452;
                                                                                                                                                                                                                                                                                                                                       rissue-Brain,
                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=21638749;    PubMed=11780052;    DOI=10.1038/414865a;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neuroprotective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=ADNP; Synonyms=KIAA0784;
Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADNP_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biol.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eutheria;
                                                                                                                                                                                                                                                                                                                                                         30-1102 FROM N.A.
                                                                                                                                 98-1102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       276:708-714(2001).
                                                                                                                                 FROM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chordata;
                                                                                                                               N.A.
                                                                                                                                                                                                                                                                                     Suyama M., Kikuno R., Miyajima N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1102 AA
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RESULT 5
Q65H2B
ID Q66HB
ID Q66HB
AC Q6DH
AC Q6DH
DT 25-C
DT 25-C
DT 25-C
DT 25-C
DT 8C-C
DT 25-C
DT 8C-C
DT 15-C
DT 1
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Best Local S
Matches
                                                                                                                                                                                                                                                                                                                       Q6DHZ8;
25-OCT-2004
25-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Transcription:
Transc
SEQUENCE FROM N.A.
TISSUE=Placenta;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SMART; SM00389; HOX; 1.
SMART; SM00355; ZNF C2H2; 8.
PROSITE; PS00027; HOMEOBOX 2; 1.
PROSITE; PS50071; HOMEOBOX 2; 1.
PROSITE; PS00028; ZINC FINGER C2H2
PROSITE; PS50157; ZINC_FINGER_C2H2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; AP250860; AAG47651.1; --
EMBL; AL034553; CAB53748.2; --
EMBL; AB018327; BAA34504.1; --
EMBL; AL080163; CAB45752.1; --
PIR; T12546; T12546.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced between the Swiss Institute of Bioinformatics and the European Bioinformatics Institute. There are a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- SUBCELLULAR LOCATION: Nuclear (Potential).
-!- TISSUE SPECIFICITY: Widely expressed. Strong expression in healing skeletal muscle, kidney and placenta. In brain, expression is stronger in the cerebrellum and cortex regions. No expression
                                                                                                                                                    Eukaryota;
Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                               Q6DHZ8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pfam; PF00046; Homeobox; 1
Pfam; PF00096; zf-C2H2; 2.
                                                                                                            NCBI_TaxID=9606;
                                                                                                                                                                                                          Homo sapiens (Human)
                                                                                                                                                                                                                                                                 Activity-dependent
                                                                                                                                                                                                                                                                                               25-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              breast cancer tissues.
-!- SIMILARITY: Contains 1 homeobox domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA-binding; Homeobox; Metal-binding; Nuclear protein; Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genew; HGNC:15766; ADNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    41 NAPVSIPQ 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAPVSIPO 361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IPR001356; Homeobox.
IPR009057; Homeodomain_like.
IPR007087; Znf_C2H2.
                                                                                                                                                 ; Metazoa;
Eutheria;
                                                                                                                                                                                                                                                                                   4 (TrEMBLrel. 28,
1 (TrEMBLrel. 28,
1 (TrEMBLrel. 28,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              74
107
165
221
221
447
489
512
622
662
754
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                                                                                                                                                                                                                                                                                                                                                                                                               PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA;
                                                                                                                                                                                                                                                                 neuroprotector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  129
129
188
244
469
510
535
                                                                                                                                                    Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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                                                                                                                                                    Craniata; Vi
Catarrhini;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 18;
Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                               1102
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                                                                                                                                                                               Vertebrata; Euteleostomi;
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(atypical)
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(atypical).
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                                                                                                                                                       Hominidae;
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L outstation -
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RESULT
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RA Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Haieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Wadin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Bana S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Pichards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J. Maley M.A.
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Best Local
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InterPro; IPR007087; Znf C2H2.

Pfam; PF00046; Homeobox; 1.

Pfam; PF00046; Homeobox; 1.

Pfam; PF00096; zf-C2H2; 1.

SMART; SM00389; HOX; 1.

SMART; SM00389; HOX; 1.

SMO0355; ZnF C2H2; 8.

PROSITE; PS0011; ABC TRANSPORTER 1; UNKNOWN 1.

PROSITE; PS50071; HOMEOBOX 2; 1.

PROSITE; PS50078; ZNC FINCER C2H2 ; 1.

PROSITE; PS50078; ZNC FINCER C2H2 ; 1.

DNA-binding; Homeobox; Nuclear protein.

SEQUENCE 1102 AA; 123446 MW; 286984E554DC766A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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GO; GO:0003700; F:transcription factor activity; IEA.
GO; GO:0008270; F:zinc ion binding; IEA.
GO; GO:0008355; P:regulation of transcription, DNA-dependent;
InterPro; IPR003439; ABC_transporter.
InterPro; IPR0031356; Homeodom.
InterPro; IPR009057; Homeodomain_like.
InterPro; IPR007087; Znf_C2H2.
                                                                                                                                                                                                                                                                                                                                         OGC9B5;
OGC7-2004 (TrEMBLrel. 28, Created)
25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation updat
25-OCT-2004 (TrEMBLrel. 28, Last annotation updat
Similar to DEHAOD167979 Debaryomyces hansenii.
ORFNames-Y4-DIO124969;
ORFNames-Y4-OTITION ORF
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Submitted (JUL-2004) to the
        Dujon
                                                                                                                                                                                                                                                       Yarrowia lipolytica CLIB99.
Eukaryota; Fungi; Ascomycota; Sac
Saccharomycetales; Dipodascaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Placenta;
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                                                                                           SEQUENCE FROM N.A.
STRAIN=CLIB99;
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                                                        Genolevures;
                                                                                                                                                                                                           NCBI_TaxID=284591;
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8; Conserv
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             Sherman
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                 Fischer
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uclear (By similarity).
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Pred. No. 18;
             G.,
                                                                                                                                                                                                                                                                                                  Saccharomycotina; Saccharomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2B6984E554DC766A CRC64;
                 Durrens
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                 Casaregola
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RA Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
RA Barnay S., Blanchin S., Beckerich J.M., Beyne E., Bleykasten C.,
RA Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
RA Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
RA Hantraye F., Hemnequin C., Jauniaux N., Joyet P., Kachouri R.,
RA Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
RA Nicaud J.M., Nikolski M., Oztas S., Ozter-Kalogeropoulos O.,
RA Pellenz S., Potier S., Richard G.F., Straub M.L., Suleau A.,
RA Swennene D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
RA Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
RA Wincker P., Souciet J.L.,
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Query Match
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Q82YW3;
Q1-JUN-2003 (TrEMBLrel. 2:
01-JUN-2003 (TrEMBLrel. 2:
01-OCT-2003 (TrEMBLrel. 2:
01-OCT-2003 (TrEMBLrel. 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FAVOR .....

STRAIN=WS83 / ATCC 700802;

MEDLINE=22550857; PubMed=12663927; DOI=10.1126/science.1080613;

MEDLINE=22550857; PubMed=12663927; DOI=10.1126/science.1080613;

Paulisen I.T., Banezjei L., Wyers G.S.A., Nelson K.E., Seshadri R.,

Paulisen I.T., Fouts D.E., Eisen J.A., Gill S.R., Heidelberg J.F.,

Read T.D., Fouts D.E., Eisen J.A., Gill S.R., Heidelberg J.F.,

Tettelin H., Dodson R.J., Umayam L.A., Brinkac L.M., Beanan M.J.,

Tettelin H., Dodson R.J., Umayam L.A., Kolonay J.F., Madupu R.,

Daugherty S.C., DeBoy R.T., Durkin S.A., Kolonay J.F., Madupu R.,

Nelson W.C., Vamathevan J.J., Tran B., Upton J., Hansen T., Shetty J.,

Nelson W.C., Vamathevan J.J., Radune D., Ketchum K.A., Dougherty B.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genoscope;
Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; CR382130; CAG80935.1; -.
SEQUENCE 445 AA; 50191 MW; DB039F405D2D9BBC CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Name=citf; OrderedLocusNames=EF3319;
Enterococcus faecalis (Streptococcus faecteria; Firmicutes; Lactobacillales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. STRAIN=CLIB99;
                                                                                                                                                                             InterPro; IPR006472; CitF.
Pfam; PF04223; CitF; 1.
PIRSE; PIRSE7009451; CittT_lyas_alpha; 1.
TIGRPAMs; TIGR01584; citF; 1.
                                                                                                                                                                                                                                                                                            TIGR, BE7319; -.

GO; GO:0009346; C:citrate lyase complex; I
GO; GO:0008814; F:citrate COA-transferase
GO; GO:0006084; P:acetyl_COA metabolism; I
                                                                                                                                                                                                                                                                                                                                                                                                                                             Enterococcus faecalis.";
Science 299:2071-2074(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Citrate lyase, alpha subunit.
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Nature 430:35-44(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fraser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=1351;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Role of mobile DNA in the evolution
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7
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                                                                                                                                                                                                                                                                                                                                                                                                              AE016957; AA082984.1;
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Similarity 6; Conserv
                                                                                                                     proteome.
470 AA;
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75.0%;
                                                                                                                        50138 MW;
                              29.8%;
75.0%;
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Last annotation update)
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   Score 36; DB Pred. No. 77; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Created)
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Pred. No.
                                                                                                                        80B3116774432C93 CRC64;
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                                                                                                                                                                                                                                                                                                                                 activity; IEA.
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                                                                Length 470;
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ralla E.,
sarbe V.,
rasten C.,
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SQUERRITA DE RECEPTOR DE LA COMPANIA DEL COM
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Q8E338;
01-MAR-2003
                                                                                                MEDLINE=21927593; PubMed=11917108; DOI=10.1073/pnas.062526099; Smoot J.C., Barbian K.D., Van Gompel J.J., Smoot L.M., Chaussee M.S. Sylva G.L., Sturdevant D.E., Ricklefs S.M., Porcella S.F., Sylva G.L., Beres S.B., Campbell D.S., Smith T.M., Zhang Q., Parkins L.D., Beres S.B., Campbell D.S., Smith T.M., Zhang Q., Kapur V., Daly J.A., Veasy L.G., Musser J.M.; "Genome sequence and comparative microarray analysis of serotype M18 group A Streptococcus strains associated with acute rheumatic fever
                                                                                                                                                                                                                                                                                                                                                                                                                                01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last samotation update)
Putative citrate lyase, alpha subunit.
Name=citF; OrderedLocusNames=spyM18 1140;
Streptcocccus pyogenes (serotype M18).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
STRAIN-MEM316 / Serotype III;
MEDLINE-22242508; PubMed=12354221;
Glaser P., Rusniok C., Buchrieser (
Glaser P., Zouine M., Couve E., Lai
                          Proc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sagalist; gbs1925; ...
InterPro; IPR006270; Strep_his_triad.
Pfam; PF04270; Strep_his_triad; 4.
TIGRFAMs; TIGR01363; strep_his_triad; 1.
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01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                             Streptococcus pyogenes
Bacteria; Firmicutes; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Streptococcus agalactiae (serotype III).
Bacteria; Firmicutes; Lactobacillales; Streptococcaceae;
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MO1. Microbiol. 45:1499-1513 (2002).

EMBL; AL766854; CAD47584.1; -.
                                                                             outbreaks.";
                                                                                                                                                                                                                                                                                     STRAIN=MGAS8232;
                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
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  L; AE010039;
GO:0009346;
                                                                                                                                                                                                                                                                                                                                                                TaxID=186103;
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                             Acad. Sci. U.S.A. 039; AAL97760.1; -
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     C:citrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     288
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             53326 MW; DFDF0453D8A929BE CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29.8%;
75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                             ; (serotype M18).
Lactobacillales; Streptococcaceae;
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Couve E., Lalioui
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     lyase
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                                                     99:4668-4673 (2002)
     complex; IEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        510
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ui L., Poyart C., Trieu-Cuot
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RESULT 11
Q8K7F1
ID Q8K7F
AC Q8K7F
DT 01-OC

Q8K7F1 PRELIMINARY; Q8K7F1; Q79X56; 01-OCT-2002 (TrEMBLrel. 22,

Created)

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RESULT
Q99ZK7
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Q99ZK7;
01-JUN-2001
                                                                      Complete
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GO; GO:0008814; F:citrate CoA-transferase activity; GO; GO:0005489; F:electron transporter activity; GO; GO:0016829; F:lyase activity; IEA. GO; GO:0006084; P:acetyl-CoA metabolism; IEA. GO; GO:0006118; P:electron transport; IEA.
                                                                                               PIRSF;
                                                                                                                                                                                                                                                                                                        01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Putative citrate lyase, alpha subunit (EC 4.1.3.6).
Name=citF; OrderedLocusNames=SPY1189;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  InterPro; IPR006472; CitF.
InterPro; IPR000049; ETF_beta.
Pfam; PF04223; CitF; 1.
PIRSF; PIRSF009451; Citrt lyas_alpha; 1.
ProDom; PD003528; ETF_beta; 1.
TIGRPAMS; TIGR01584; CitF; 1.
                                                                                                                                                                                                                                                                     Streptococcus.
NCBI_TaxID=1314;
                                                                                      PIRSF; PIRSF009451; Citrt_lyas_alpha; 1.
TIGRFAMs; TIGR01584; citF; 1.
                                                                                                                                                                                                                                                                                          Bacteria; Firmicutes; Lactobacillales; Streptococcaceae;
                                                                                                                                                                                                                                                                                                  Streptococcus pyogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Complete proteome; Lyase.
SEQUENCE 510 AA; 54918 MW;
                                                                                                        InterPro; IPR006472; CitF.
Pfam; PF04223; CitF; 1.
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215
                 41 NAPVSIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                               6; Conserv
                                  Similarity 6; Conserv
NTPISIPO 222
                                                                      proteome;
510 AA;
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                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            F:citrate CoA-transferase activity; IEA.
                 48
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                                                                     Lyase.
54990 MW;
                                           29.8%;
75.0%;
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                                             Pred.
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                                                                      459833B693E2EFA7 CRC64;
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                                   Mismatches
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                                             36;
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                                                                                                                                                                                                                                                                                                                                                                 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                         84;
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REMBL; AP005144; BAC641291; ...

ROGO:0009346; C:citrate lyase complex; IEA.

ROGO:0008814; F:citrate CoA-transferase activity; IEA.

ROGO:0005489; F:electron transporter activity; IEA.

ROGO:00016829; F:lyase activity; IEA.

ROGO:0006181; P:acetyl-CoA metabolism; IEA.

ROGO:0006118; P:electron transport; IEA.

ROGO:000618; P:electron tra
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SETRAIX=MGAS315 / Serotype M3;

MEDLINE=22133808; PubMed=12122206; DOI=10.1073/pnas.152298499;

Beres S.B., Sylva G.L., Barbian K.D., Lei B., Hoff J.S.,

Beres S.B., Sylva G.L., Barbian K.D., Lei B., Hoff J.S.,

Campbell D.S., Smith T.M., McCormick J.K., Leung D.Y.M.,

Schlievert P.M., Musser J.M.;

"Genome sequence of a serotype M3 strain of group A Streptococy

phage-encoded toxins, the high-virulence phenotype, and clone
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25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Putative citrate lyase, alpha subunit.
Name=citf; OrderedLocusNames=SPs1034, SpyM3_0834;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nakagawa I., Kurokawa K., Yamashita A., Nakata M., Okahashi N., Kawabata S., Yamazaki K., Shiba T., Y Hayashi H., Hattori M., Hamada S.; "Genome sequence of an M3 strain of Streptococcus large-scale genomic rearrangement in invasive strainto phage evolution.";
Genome Res. 13:1042-1055(2003).
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                                                                                                                                                                                                                                                                                                      01-MAR-2003 (TrEMBLrel 23, Created)
01-MAR-2003 (TrEMBLrel 25, Last sequence update)
01-CCT-2003 (TrEMBLrel 25, Last annotation updat)
Putative citrate lyase, alfa subunit (EC 4.1.3.6)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=22663278; PubMed=12799345; DOI=10.1101/gr.1096703;
Nakagawa I., Kurokawa K., Yamashita A., Nakata M., Tomiyasu Y.,
Okahashi N., Kawabata S., Yamazaki K., Shiba T., Yasunaga T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=198466;
SEQUENCE FROM N.A.
STRAIN=UA159 / ATCC 700610 /
                                                                                                                                                                                                                                                   Putative citrate lyase, alfa subunit (Name=cilA; OrderedLocusNames=SMU.1021;
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                                                                                                      NCBI_TaxID=1309;
                                                                                                                                                     Streptococcus
                                                                                                                                                                                            Bacteria; Firmicutes;
                                                                                                                                                                                                                                Streptococcus mutans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q8DUC1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       QBDUC1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
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Lactobacillales; Streptococcaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29.8%;
                                                                                                                                                                                            Lactobacillales;
   Serotype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               F84036E9F2B0D9A8 CRC64;
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   O
                                                                                                                                                                                        Streptococcaceae;
                                                                                                                                                                                                                                                                                              tion update)
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ains and new insights
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RESULT 13
Q912G
AC Q912G
DT 01-M
DT 01
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Best Local Similarity
Matches 6; Conser
                                                                                                                                                                         Query Match
Best Local
                                                                                                                                       Matches
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Q912G3;
01-MAR-2001
01-MAR-2001
01-MAR-2004
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Pfam; PF04223; CitF; 1.

PIRSF; PIRSF009451; Citrt_lyas_alpha;

TIGRFAMs; TIGR01584; citF; 1.

Complete proteome.

SEQUENCE 511 AA; 55455 MW; FF966EE
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EMBL, AE014941; AANS9721.1; -
CO; GC:0009346; C:citrate lyase complex; IEA.
GO; GO:0008815; F:citrate (pro-3S)-lyase activity; I
GO; GO:0008814; F:citrate CoA-transferase activity; GO; GO:0006814; P:acetyl-CoA metabolism; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.

STRAIN=ATCC 15692 / PAO1;

STRAIN=ATCC 15692 / PAO1;

STRAIN=20437337; PubMed=10984043; DOI=10.1038/35023079;

MEDLINB=20437337; PubMed=10984043; DOI=10.1038/35023079;

MEDLINB=20437337; PubMed=10984043; DOI=10.1038/35023079;

MEDLINB=20437337; PubMed=10984043; DOI=10.1038/35023079;

MEDLINB=20437337; PubMed=10.1038/35023079;

MEDLINB=20437337; PubMed=10.1038/35023079;

MEDLINB=20437337; PubMed=10.1038/35023079;

MEDLINB=20437337; PubMed=10.1038/35023079;

MEDLINB=20437337; PubMed=10.1038/35023079;

MEDLINB=20437337; PubMed=10.984043; DOI=10.1038/35023079;

MEDLINB=2043737; PubMed=10.984043; DOI=10.1038/35023079;

MEDLINB=20437337; PubMed=10.984043; DOI=10.1038/35023079;

MEDLINB=2044744, MEDLINB=204044, MEDLINB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pathogen.";
                                                                                                                                                                                                                                                                                   InterPro; IPR000345; CytC_heme_BS.
InterPro; IPR009056; CytoChrome_C:
PROSITE; PS00190; CYTOCHROME_C; UNKNOWN_2.
Complete proteome; Hypothetical protein.
SEQUENCE 631 AA; 70904 MW; E72A4692AEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Complete genome sequence of Pseudomonas aeruginosa PAO1, opportunistic pathogen.";
Nature 406:959-964 (2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical protein.
OrderedLocusNames=PA1941;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Genome sequence of Streptococcus mutans UA159, a cariogenic dental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=22295063; PubMed=12397186; DOI=10.1073/pnas.172501299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AE004620; AAG05329.1; -. PIR; B83404; B83404.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=287;
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263
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                                                                                                                                       Similarity
7; Conserv
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NAPVSYPQ
                                                                   NAPVSIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                           Conservative
270
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87.5%;
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75.0%;
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Last annotation updat
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                                                                                                                                                                             Score 36;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 36; DB
Pred. No. 84;
                                                                                                                                                                                                                                                                                       l protein.
E72A4692AEEB75F1 CRC64;
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                                                                                                                                               Mismatches
                                                                                                                                                                                 1.1e+02;
                                                                                                                                                                                                                   DB 2;
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                                                                                                                                                                                                                       Length 631;
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in S.P.,
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Best Local S
Matches 6
SEQUENCE FROM N.A.

STRAIN=C57BL/6J;

MEDLINE=2354683; PubMed=12466851; DOI=10.1038/nature01266;

MEDLINE=2354683; PubMed=12466851; DOI=10.1038/nature01266;

MEDLINE=2354683; PubMed=12466851; DOI=10.1038/nature01266;

MEDLINE=2354683; PubMed=12466851; DOI=10.1038/nature01266;

MEDLINE=2354683; PubMed=1., Kondo S.,

Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,

Yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,

Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,

Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,

Schriml L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W.,

Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,

Dalla E., Dragani T.A., Fletcher C.F., Forrest A., Frazer K.S.,

Dalla E., Dragani T.A., Fletcher C.F., Forrest A., Frazer K.S.,

Captariand T., Gariboldi M., Gissi C., Godzik A., Gough J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q8BX17;
10-OCT-2003
10-OCT-2003
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ProDom; PD000018; WD40; 1.

SMART; SM00320; WD40; 11.

PROSITE; PS00589; PTS HPR SER; UNKNOWN 1.

PROSITE; PS00678; WD REPEATS 1; UNKNOWN 2.

PROSITE; PS50082; WD REPEATS 2; 2.

PROSITE; PS50294; WD REPEATS REGION; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA Res. 11:167-180(2004).

- SIMILARITY: Contains 9 WD repeats.
EMBL, AK131137; BAD21387.1; -
InterPro; IPR002114; HPT SerP S.
InterPro; IPR001680; WD40.
InterPro; IPR001046; WD40.
InterPro; IPR001046; WD40.
Pfam; PF00400; WD40; 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Okazaki N., Kikuno R., Ohara R., Inamoto S., Koseki H., Hiraoka S., Saga Y., Kitamura H., Nakagawa T., Nagase T., Ohara O., Koga H.; "Prediction of the Coding Sequences of Mouse Homologues of FLJ Genes: The Complete Nucleotide Sequences of 110 Mouse FLJ-Homologous cDNAs Identified by Screening of Terminal Sequences of cDNA Clones Randomly Sampled from Size-Fractionated Libraries.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JUL-2004
MFLJ00137 p
                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-JUL-2004
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=Gemin5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Repeat; WD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (Rel. 42, Created)
(Rel. 42, Last sequence update)
(Rel. 44, Last annotation update)
ed protein 5 (Gemin5).
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IBLrel. 27,
IBLrel. 27,
I (Fragment)
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Rodentia;
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Rodentia;
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75.0%;
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Last sequence update)
Last annotation update)
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2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 36; DB 2;
Pred. No. 2.5e+0;
                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1502 AA.
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RA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
RA Magshima T., Numata K., Okido T., Pavan W.J., Pertea G., pesole G.,
RA Petrovsky N., Pillai R., Pontius J.U., Oi D., Ramachandran S.,
RA Petrovsky N., Pillai R., Pontius J.U., Oi D., Ramachandran S.,
RA Ravasi T., Reed J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M.,
RA Sandelin A., Schneider C., Semple C.A., Setou M., Shimada K.,
RA Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M.,
Verardo R., Wagner L., Wahlestedt C., Wang Y., Watanabe Y., Wells C.,
RA Vilming L.G., Wynnhaw-Boris A., Yanagisawa M., Yang I., Yang L.,
RA Wilming L.G., Zavolan M., Zhu Y., Zimmer A., Carninci P., Hayatsu N.,
RA Yuan Z., Zavolan M., Zhu Y., Zimmer A., Carninci P., Hayatsu N.,
RA Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K.,
RA Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K.,
RA Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K.,
RA Hara A., Hashizume W., Imotani K., Ishii Y., Itoh M., Kagawa I.,
RA Miyazaki A., Sakai K., Sasaki D., Shibata K., Shinagawa A.,
Yasunishi A., Yoshino M., Waterston R., Lander E.S., Rogers J.,
RA Birney E., Hayashizaki Y.,
RA Birney E., Hayashizaki Y.,
Nature 420.563-571(700)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nature 420:563-573(2002).

Nature 420:563-573(2002).

Nature 420:563-573(2002).

It is played an essential role in spliceosomal splicing in the cytoplasm and is required for pre-mRNA splicing in the nucleus.

It subunit: Part of the core SMN complex that contains SMN1, SIP1, GRMINIZ: GEMIN3, GEMIN4, GEMIN5, GEMIN6 and GEMIN7: Interacts GRMIN2: GEMIN3, GEMIN4, GEMIN5, GEMIN6 and GEMIN7: Interacts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a celebrate of the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AK049216; BAC33614.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entities requires a
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Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., Kin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -!- SIMILARITY: Contains 13 WD repeats.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in nuclear bodies called gems (Gemini of Caji often in proximity to Cajal (coiled) bodies. cytoplasm (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUBCELLULAR LOCATION: Nuclear,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         directly with SMN1, SNRPB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                license agreement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNRPD1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  found both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (See http://www.isb-sib.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNRPD2,
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Cajal bodies) that are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GEMIN7. Interacts
SNRPD3 and SNRPE (By
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             found in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           restrictions
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MGD; MGI:2449311; Gemin5.

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REPEAT
REPEAT
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DOMAIN
DOMAIN
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REPEAT
                                                                                                                                                                                         SMART; SM00320; WD40; 13.

PROSITE; PS0062; WD REPEATS 1; 3.

PROSITE; PS50082; WD REPEATS 2; 3.

PROSITE; PS50294; WD REPEATS REGION; 1.
                                                                                                                                                                     Spliceosome;
                                                                                                                                                                                                                                   PRINTS; PR00320; GPROTEINBRPT
ProDom; PD000018; WD40; 1.
                                                                                                                                                                                                                                                                     InterPro;
                                                                                                                                                                                                                                                                                         InterPro;
                                                                                                                                                                                                                                                          PF00400; WD40;
                                                                                                                                                                                 coil
                                                                                                                                                                                                                                                                 IPR011048; Cyt_cd1_haem_C.
IPR008941; TPR-like.
IPR001680; WD40.
                                                                                           WD
107
150
193
280
333
                                                                                                                                                            repeat.
AA;
                                                                                                                                                                               processing;
 166562
      WD 1.
WD 2.
WD 3.
WD 4.
WD 5.
WD 6.
WD 7.
WD 9.
WD 10.
WD 11.
WD 13.
D 13.
Poly-Lys.
Coiled coil (Potential).
 ₹.
                                                                                                                                                                                 mRNA
46F98CCCE70E2971 CRC64;
                                                                                                                                                                               splicing; Nuclear protein; Repeat;
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Page 8
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Query Match 29.8%; Score 36; DB 1; Length 1502;
Best Local Similarity 75.0%; Pred. No. 2.7e+02;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 41 NAPVSIPQ 48
|\| | | | | | |
Db 1420 NAPVSLPE 1427

Search completed: February 23, 2005, 19:34:18
Job time: 116.35 secs
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Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                             OM protein -
                                                           Database :
                                                                                      Post-processing: Minimum Match 0%
Maximum Match 10
Listing first 45
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Maximum DB
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Perfect score:
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                                                                                                                                                 seq length: 0
seq length: 2000000000
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                                                                                                                                                                                                                                                         BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                        US-09-267-511-4
121
PIR 79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
                                                                                                                                                                                                                            283416 seqs, 96216763 residues
                                                                                                                                                                                                                                                                                                                                                                                               February 23,
                                                                                                                                                                                                                                                                                                        Copyright
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                                                                                                                                                                                                                                                                                                                                                               2005, 19:22:32; Search time 24.3616 Seconds (without alignments) 347.559 Million cell updates/sec
                                                                                        100%
45 ฮเ
                                                                                         summaries
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Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Conservative

0

1;

Indels

0

Gaps

0

ALIGNMENTS

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A;Molecule type: DNA
A;Residues: 1-631 <STO>
A;Cross-references: UNIPROT:0912G3; GB:AE004620; GB:AE004091; NID:g9947929; PIDN:AAG053.
A;Experimental source: strain PAO1
C;Genetics:
                                                                                                                                                                                                                                                                              C;Accession: B83404

R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; B adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim ; Lory, S.; Olson, M.V.

Nature 406, 959-964, 2000

A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic pathu A;Reference number: A82950; MUID:20437337; PMID:10984043

A;Accession: B83404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypothetical protein DKFZp586K2120.1 - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 23-Jul-1999 #sequence_revision 23-Jul-1999 #text_change 09-Jul-2004
C;Accession: T12546
R;Wambutt, R; Heubner, D.; Mewes, H.W.; Gassenhuber, J.; Wiemann, S.
submitted to the protein Sequence Database, June 1999
A;Reference number: Z17524
A;Accession: T12546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypothetical protein PA1941 [imported] - Pseudomonas aeruginosa (strain PAO1) C;Species: Pseudomonas aeruginosa C;Species: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
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A;Residues: 1-1005 <MAM>
A;Residues: 1-1005 <MAM>
A;Cross-references: UNIPROT:Q9H2P0; EMBL:AL080163
A;Experimental source: adult uterus; clone DKFZp586K2120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Status: preliminary
Query Match
Best Local Similarity
Matches 7; Conserv
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Best Local S
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                                                                                                          PA1941
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 100.0%; Score 41; DB 2; Similarity 100.0%; Pred. No. 3.2; 8; Conservative 0. wirman-
                          29.8%;
     Score 36; DB
Pred. No. 20;
0; Mismatches
                                  20;
                                                       2
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                                                     Length 631
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hypothetical protein SMa2233 [imported] - Sinorhizobium meliloti (strain 1021) magaplasm C;Species: Sinorhizobium meliloti (c;Species: Sinorhizobium meliloti (c;Species: Sinorhizobium meliloti (c;Date: 24-Aug-2001 #sequence_revision 24-Aug-2001 #text_change 09-Jul-2004 (c;Date: 24-Aug-2001 #sequence, T:; Komp, C.; Abola, A.P.; Barloy-Hubler, F.; Bows R;Barnett, M.J.; Fisher, R.F.; Jones, T.; Komp, C.; Abola, A.P.; Barloy-Hubler, F.; Bows R; Kalman, S.; Keating, D.H.; Palm, C.; Peck, M.C.; Surzycki, R.; Wells, D.H.; Yeh, K.C. Proc. Natl. Acad. Sci. U.S.A. 98, 9883-9888, 2001
A;Title: Nucleotide sequence and predicted functions of the entire Sinorhizobium melilot A;Reference number: A95262; MUID:21396509; PMID:11481432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Residues: 1-299 <KUR>
A;Cross-references: UNIFROT;O92XPO; GB:AB006469; PIDN:AAK65862.1; PID:g14524369; GSPDB: A;Cross-references: UNIFROT;O92XPO; megaplasmid pSymA
A;Experimental source: strain 1021, megaplasmid pSymA
R;Galibert, F; Finan, T.M.; Long, S.R.; Puhler, A.; Abola, P.; Ampe, F.; Barloy-Hubler
pela, D.; Chain, P.; Cowie, A.; Davis, R.W.; Dreano, S.; Federspiel, N.A.; Fisher, R.F.
L; Hyman, R.W.; Jones, T.
Science 293, 668-672, 2001
A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure
h;Authors: Kahn, D.; Kahn, M.; Vorholter, F.J.; Weidner, S.; Wells, D.H.; Wong, K.; Yeh, K
A;Reference number: A96039; MUID:21368234; PMID:11474104
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                                                                                                                                                                                                                                                                                                                       R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dods.; Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E. Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L. Nature 390, 364-370, 1997
A;Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, Smith, H.O.; Woese, C.R.; Venter, J.C.
A;Title: The complete genome sequence of the hyperthermophilic, sulfate-reducing archa A;Reference number: A69250; MUID:98049343; PMID:9389475
A;Accession: G69440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  conserved hypothetical protein AF1528 - Archaeoglobus fulgidus
C;Species: Archaeoglobus fulgidus
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Status: preliminary A;Molecule type: DNA
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                                                                                                                                                                                                                          A;Residues: 1-136 <KLE>
A;Cross-references: UNIPROT:O28744; GB:AE000997; GB:AE000782; NID:g2689320;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Accession: G69440
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Best Local
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                                                                                                                                                                                                                                                     ;Molecule type: DNA
;Residues: 1-136 <KLE>
                                                                                                                                                                                                                                                                                                 Status: preliminary; nucleic
                                                                                                                                                                      Query Match
                                                                                                                                              Local
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                29
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Similarity 100.0%;
7; Conservative
                                                                                                                                              Similarity
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                                                                    NAPVSIPO 48
                                                                                                                     Conservative
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                   36
                                                                                                                                              28.1%;
                                                                                                                                                                                                                                                                                                       acid
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Pred. No.
                                                                                                                        Score 34; DB:
Pred. No. 9.3;
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 red. No. 14;
Mismatches
                                                                                                                                                                                                                                                                                                    sequence
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                                                                                                                                                                            Length 136;
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                                                                                                                                                                                                                                                                                                         translation not
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                                AH0813
probable ethanolamine utilization protein EutA eutA [imported]
C;Species; Salmonella enterica subsp. enterica serovar Typhi
A;Note: this species has also been called Salmonella typhi
C;Date: 09-Nov-2001 #sequence_revision 09-Nov-2001 #text_change
C;Accession: AH0813
C;Accession: AH0813
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R; Parkhill, J.; Dougan, G.; James, th, T.; Connerton, P.; Cronin, A.;
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NAPVATPO

O'Gaora,

K.D.; Thomson, Davis, P.; Dav

Davies,

N.R.; Pickard, D.; Wain, J.; ies, R.M.; Dowd, L.; White, N

Churcher,

#text_change

18-Nov-2002

Salmonella

enterica

guit

41 NAPVSIPQ 48

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N;Alternate names: fmr2 protein
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text_chan
C;Accession: T30248
R;Chakrabarti, L.; Bristulf, J.; Foss, G.S.; Davies, K.E.
Hum. Mol. Genet. 7, 441-448, 1998
A;Title: Expression of the murine homologue of FMR2 in mouse
A;Title: Expression of the murine homologue of FMR2 in mouse
A;Reference number: Z20786; MUID:98133924; PMID:9467002
A;Accession: T30248
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A;Residues: 1-1272 <CHA>
A;Cross-references: UNIPROT:055112;
                                                                                                                                                                                                                                                                                                                                                                                                               th, T.; Connerton, P.; Cronin, A.; Davis, P.; Davies, R.M.; Dow, S.; Moule, S.; O'Gaora, P.
Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.;
A;Title: Complete genome sequence of a multiple drug resistant A;Reference number: AB0502; MUID:21534947; PMID:11677608
A;Accession: AI0632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  trp repressor binding protein [imported] - Salmonella enterica subsp. enterica serovar T C;Species: Salmonella enterica subsp. enterica serovar Typhi A;Note: this species has also been called Salmonella typhi C;Date: 09-Nov-2001 #sequence_revision 09-Nov-2001 #text_change 18-Nov-2002 C;Accession: Al632 C;Accession: Al632 C;Accession: Al632 C;Accession: Al632 C;Accession: Al632 C;Accession: Al632 C;Caccession: Al632 C;Accession: Al632 C;Caccession: Al632 C;Accession: Al632 C;Caccession: Al632 C;Caccess
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AI0632
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Status: preliminary; translated from
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                                                                                                                                                                                                                                                                        A;Molecule type: DNA
A;Residues: 1-198 <PAR>
A;Cross-references: GB:
                                                                                                                                    A;Gene: STY1155
C;Superfamily: trp repressor-binding
C;Keywords: flavoprotein
                                                                                                                                                                                                                                                                                                                                                                                  A; Status: preliminary
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Best Local S
Matches 6
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Similarity 6; Conserv
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   Conservative
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75.0%;
                                        27.3%;
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Pred. No. 1.26
1; Mismatches
       Score 33; DB
Pred. No. 23;
1; Mismatches
                                                                                                                                                                                                                                                                                   PIDN:CAD08244.1; PID:g16502291; GSPDB:GN00176
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                                                                                                                                                                            protein;
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brain

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C.Accession: S47896
C.Accession: S47896
R;Kamdar, K.P.; Shelton, M.E.; Finnerty, V.
Renetics 137, 791-801, 1994
A;Title: The Drosophila molybdenum cofactor gene cinnamon
A;Title: The Drosophila molybdenum cofactor gene cinnamon
A;Title: The Drosophila molybdenum cofactor gene cinnamon
                A; Molecule type: mRNA
A; Residues: 1-601 <KAM>
A; Cross-references: EMBL:L19876; NID:g797288; PIDN:AAA65877.1;
A; Experimental source: developmental stage embryo
C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso, Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K.; ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.; Kim, C.C.A.; Li, J.H.; Li, Y.; Lin, S.X.; Liu, S.A.; Luros, J.S.; Maiti, R.; Marziali, Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallon, Ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A;Accession: H86465
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                                                                                                                                                                                                                                            probable molybdopterin biosynthesis protein cinnamon - fruit fly C;Species: Drosophila melanogaster C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change C;Accession: S47896
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A;Molecule type: DNA
A;Residues: 1-479 <STO>
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C;Superfamily: Escherichia coli hypothetical protein b2451
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A; Residues: 1-467 < PAR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
C;Accession: H86465
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C;Species: Arabidopsis thaliana (mouse-ear cress)
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Genetics:
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Best Local Similarity
Matches 6; Conserv
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Best Local
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Pred. No. 63;
4; Mismatches
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Pred. No.
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61;
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Salmonella enterica s
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ovo protein - fruit fly (Drosophila melanogaster)
C;Species: Drosophila melanogaster
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995
C;Accession: S16356
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A; Molecule type: DNA
A; Residues: 1-1119 <SCH>
A; Cross-references: UNIPROT: Q9P3N5; EMBL: AL389891; GSPDB: GN00116; NCSP: B7F18.140
A; Cross-references: BAC clone B7F18; strain OR74A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Date: Z1-UL T50995
C;Accession: T50995
U.; Aign
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C;Function:
C;Function:
A;Pathway: molybdopterin biosynthesis
C;Superfamily: bifunctional molybdenum cofactor biosynthesis pro
C;Keywords: molybdopterin biosynthesis; multifunctional enzyme;
                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-1213 <MEV>
A; Cross-references: UNIP
                                                                                                                                                                                                                          EMBO J. 10, 2259-2266, 1991

A;Title: The ovo gene of Drosophila encodes a zinc finger protein A;Reference number: S16356; MUID:91293102; PMID:1712294

A;Accession: S16356
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Neurospora crassa
C;Date: 21-Jul-2000 #sequence_revision 21-Jul-2000 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                            R; Mevel-Ninio,
                                                                                                                                                                                                                                                                                                                                                                              S16356
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                                                                                         ;Gene: FlyBase:ovo
;Cross-references: FlyBase:FBgn0003028
;Introns: 931/3; 1152/3
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les 6; Conserv
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                                                  Similarity
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NAPVSIPO 48
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71.4%;
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1.6e+02;
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                                             DB 2; 1
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hypothetical protein T05A10.1 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
C;Accession: T24490
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Rischmitted to the EMBL Data Library, November 1995
A;Reference number: Z19898
A;Accession: T24490
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C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
C;Accession: T25754
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                                                                                       hypothetical protein al10976 [imported] - Nostoc sp. (strain PCC 7120)

C;Species: Nostoc sp. PCC 7120

A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. strain PCC 7120

C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change 09-Jul-2004

C;Date: 14-Dec-2011 #sequence_revision 14-Dec-2001 #text_change 09-Jul-2004

C;Accession: AE1928

R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, A.;
Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Takazawa, M.; Yamada, M.; Yasuda, M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-97 <WIL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     submitted to the EMBL Data Library, September 1996
A; Description: The sequence of C. elegans cosmid F45E4
A; Reference number: Z20082
A; Accession: T25754
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A;Introns: 188/3; 240/3; 420/1; 570/3; 596/1; 732/3; 778/3; 851/3; 1359/2; 1394/2;
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A;Molecule type: DNA
DNA Res. 8, 205-213, 2001

A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium Ana A;Reference number: AB1807; MUID:21595285; PMID:11759840

A;Accession: AE1928
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Pred. No. 3.6e
1; Mismatches
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Pred. No. 16;
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Search completed: February 23, Job time : 25.3616 secs

2005, 19:35:04

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RESULT 15
AE0243
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A;Residues: 1-249 <KDS;
A;Cross-references: UNIPROT:Q8YY75; GB:BA000019; PIDN:BAB72933.1; PID:g17130322; GSPDB:G
A;Experimental source: strain PCC 7120
C;Genetics:
                                                                                                                                                                                                                                 A;Title: Genome sequence of Yersinia pestis, the causative agent of plague. A;Reference number: AB0001; MUID:21470413; PMID:11586360 A;Accession: AB0243
                                                                                                                                                                                                                                                                                   R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, M.B. deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, G.; il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barrell, Nature 413, 523-527, 2001
                                                                                                                                                                                                                                                                                                                                                   hypothetical protein YPO1996 [imported] - Yersinia pestis (strain CO92) C;Species: Yersinia pestis C;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 09-Jul-2004 C;Accession: AE0243
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A;Gene: YPO1996
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A;Residues: 1-301 <KUR>
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Minimum
Maximum
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Perfect score:
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ALIGNMENTS

RESULT 1 AAW64676 Human ADNF-III derived active peptide. 04-NOV-1998 AAW64676; AAW64676 standard; protein; (first entry) 9 ₽

HIV infection Activity dependent neurotrophic factor III: ADNF-III; ADNP; cell death; activity dependent neuroprotective protein; neurone; excito-toxicity; spinal cord; hippocampus; cerebral cortex; cholinergic; beta-amyloid; nethyl-D-aspartate; Alzheimer's disease; human immunodeficiency virus;

Homo sapiens.

WO9835042-A2

13-AUG-1998.

06-FEB-1998; 98WO-US002485.

07-FEB-1997; (USSH) US SEC HEALTH & HUMAN SERVICES. 97US-0037404P

Gozes I, Brenneman DE, Bassan M;

WPI; 1998-447239/38.

Activity dependent neurotrophic factor III polypeptide - useful therapeutically to prevent neuronal cell death associated with e.g. HIV infection, excito-toxicity or Alzheimer's disease.

Example 4c; Page 67; 121pp; English.

This sequence represents a peptide used in a method which isolates a novel activity dependent neurotrophic factor III, ADMF-III (also known as activity dependent neuroprotective protein, ADMF). ADMF-III polypeptides can be used to prevent neuronal cell death, of e.g. the spinal cord, hippocampal, cerebral cortical or cholinergic neurones associated with e.g. HIV infection, excito-toxicity induced by N-methyl-D-aspartate stimulation or beta-amyloid peptide in Alzheimer's disease. The polypeptides can also be combined with a carrier to alleviate learning impairment produced by cholingeric blockage in Alzheimer's patients. The nucleic acids are useful in polypeptide production and to detect ADNF III

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                         The present sequence is the human Activity Dependent Neurotrophic Factor (ADNF) -9 active peptide sequence. The ADNF III-8 or NAP peptide was synthesised based on the structural homology to ADNF-9 peptide and hsp60. It is also used for immunological detection of cloned ADNF III. ADNF III is also called an Activity Dependent Neuroprotective Protein (ADNF). The human gene was mapped to chromosome 20413.2 and is linked to autosomal dominant nocturnal frontal-lobe epilepsy (ADNFLE) gene. It is expressed in the astrocytes, brain and also in foetal lung and endocrine tissues. This sequence has homology to ADNF I and hsp60, heat shock protein and PIF1, a DNA repair protein. The ADNF III polypeptides are useful for the treatment of neurological deficiencies and for prevention of neuronal cell death associated with gp120, the envelope protein from HIV, N-methyl D-Aspartic acid (excito-toxicity); tetrodotoxin (blockage of electrical
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                                                                                                    The present invention relates to the treatment of a condition associated with foetal alcohol syndrome (FAS), invig administering an activity dependent neurotropic factor (ADNF). ADNFs of the present invention may also be used to treat neurological deficiencies and prevent neuronal cel death. The present sequence is an ADNF peptide
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J. 1.8e+06;
O;
                                        DB 3; L
                                                   Length
                                                                                                                                                                                                                                                                                                Zamostiano
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                             Indels
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RESULT 5
AAB72315
ID AAB72
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AAB7237
XX
AC AAB7
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DT 16-M
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DE Acti
XX
KW Acti
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KW Acti
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Neu
XW vasc
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                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                     The present invention relates to the treatment of a condition associated with foetal alcohol syndrome (FAS), involving administering an activity dependent neurotropic factor (ADNF), ADNFs of the present invention may also be used to treat neurological deficiencies and prevent neuronal cell death. The present sequence is an ADNF peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treating condition associated with fetal alcohol syndrome in a subject exposed to alcohol in utero or reducing neuronal death, involves administering activity dependent neurotrophic factors I and/or III.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYRA-)
(USSH )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Activity dependent neurotrophic factor; ADNF; FAS; foetal alcohol syndrome; gene therapy; neurological deficiency; neuronal cell death.
                                                        Activity dependent neurotrophic factor I; neuronal cell death; Alzheimer's disease; vasoactive intestinal peptide; foetal deat
             WO200112654-A2
                                   Synthetic.
                                                                                                       Activity dependent neurotrophic factor I (ADNF I)
                                                                                                                              16-MAY-2001
                                                                                                                                                                             AAB72315
                                                                                                                                                                                                                                                                                                                                  Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Brenneman DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
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                                                                                                                                                                                                                                                                                Local Similarity
nes 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000-601940/57
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                                                                                                                                                                                                                                     SALLRSIPA
                                                                                                                                                                             standard;
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                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                             (first entry)
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                                                                                                                                                                            peptide; 9
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                                                                                                                                                                                                                                                                                              33.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gozes I, Bassan
                                                                                                                                                                                                                                                                                              Score 40; pred. No.
                                                                                                                                                                                                                                                                                   <u>.</u>
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                                                                                                                                                                                                                                                                                   Mismatches
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                                                       r I; ADNF I; ADNF III; body weight; ase; oxidative stress; VIP; death; foetal alcohol syndrome.
                                                                                                                                                                                                                                                                                  . 1.8e+06;
ches 0;
                                                                                                                                                                                                                                                                                                         DB 3;
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                                                                                                                                                                                                                                                                                                         Length 9;
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(ADNF I) or ADNF III polypeptide. Sequences AAB72315 - AAB72321 represent
CC ADNF II active core site peptides. Sequences AAB72322 - AAB72326 represent
CC ADNF III active core site peptides. Sequences AAB72322 - AAB72326 represent
CC composition containing either ADNF I or ADNF III and a pharmaceutical
CC neuronal cell death, e.g. death of spinal cord neurons, hippocampal
CC neurons, cerebral cortical neurons and cholinergic neurons, in a patient
CC infected with a virus, e.g. human immunodeficiency virus (HIV). The
CC neuronal cell death is associated with excito-toxicity induced by N-
CC neuronal cell death is associated with excito-toxicity induced by N-
CC neuronal cell death is associated with excito-toxicity induced by N-
CC neuronal cell death is associated with excito-toxicity induced by beta-amyloid
CC peptide in an Alzheimer's disease patient, or induced by beta-amyloid
CC peptide in an Alzheimer's disease patient, or induced by cholinergic
CC blockade. ADNF I, ADNF III and the pharmaceutical composition are also
CC useful for treating oxidative stress in a patient, for reducing a
CC condition, such as decreased body weight, decreased brain weight,
CC decreased level of vasoactive intestinal peptide (VIP) mRNA, and foetal
CC death, associated with foetal alcohol syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel Activity Dependent Neurotrophic Factor I oxidative stress, reducing neuronal cell death associated with fetal alcohol syndrome.
31-MAY-2000; 2000US-0208944P
08-FEB-2001; 2001US-0267805P
                                                                                                                                                                                        ADNP; Activity Dependent Neurotrophic Factor; nootropic; neuroprotective; cerebroprotective; antidiabetic; neuroleptic; anticonvulsant; anti-HIV; antiparkinsonian; tranquilizer; antialcoholic; vulnerary; antibacterial; antiinflammatory; antidote; ophthalmological; muscular; vasodilator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-AUG-2000;
                                              31-MAY-2001; 2001WO-US017758
                                                                                                             WO200192333-A2
                                                                                                                                           Synthetic
                                                                                                                                                                         NMDA receptor.
                                                                                                                                                                                                                                                                        ADNF I
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(USSH )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41
                                                                                                                                                                                                                                                                       polypeptide active core site peptide sequence
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                                                                                                                                                                                                                                                                                                                                                                      standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                  peptide; 9
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Pred. No.
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RESULT 7
ABR39741
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Best Local Similarity
Watches 9; Conservi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC parkinson's disease, progressive supranuclear palay, routecters by proceed to a trophy, clivo-pontocerebellar atrophy, parameoplastic cerebellar atrophy, parameoplastic cerebellar atrophy, parameoplastic cerebellar controlled as Friedreich's ataxia, diabetes, peripheral neuropathy, retinal neuronal concerebellar controlled systems such as cerebral concerebellar concerebellar controlled systems such as cerebral concerebellar cerebral concerebellar concerebel
     ZZXBXBXBXB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            subject that involves administering pre- or postnatally an Activity pependent Neurotrophic Factor (ADNF) to the subject. The ADNF polypeptide is an ADNF I and/or an ADNF III polypeptide comprising the core active site sequences ABB07215 or ABB07216. The method is useful for improving learning and/or memory in a subject; for treating a normal or old subject afflicted with neuropathology, Alzheimer's disease, Down's syndrome, normal mental capacity, mental retardation, for the treatment of central motor systems including degenerative conditions affecting the basal ganglia (such as Huntington's disease, Wilson's disease, striatonigral ganglia (such as Huntington's disease, wilson's disease, Striatonigral degeneration, corticobasal ganglionic degeneration). Tourette's syndrome,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention provides a method of improving learning and/or memory in a subject that involves administering pre- or postnatally an Activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 51; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Spong CY, Brenneman
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(USSH) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hyperglycemia, hy present sequence polypeptide
ADNF I; Activity Dependent Neurotrophic Factor I; neuroprotective; anti-HIV; neuroleptic; antiparkinsonian; nootropic; ADNF-9; SAL.
                                                                           ADNF I active core peptide fragment SAL.
                                                                                                                                  23-JUN-2003
                                                                                                                                                                                                                                      ABR39741 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                    41
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                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       represents the active core site sequence of the ADNF
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 40; DB; Pred. No. 1.8
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RESULT 8
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Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to Activity Dependent Neurotrophic Factor I (ADNF I) complex polypeptide selected from sequences ABR39744-754. The ADNF I complex polypeptide is useful for reducing neuronal cell death in conditions such as HIV infection, treating oxidative strees in a patient, reducing a condition associated with fetal alcohol syndrome, or improving learning and/or memory in a subject with e.g. Alzheimer's disease or Down's syndrome. The ADNF complex polypeptides are also useful in designing a drug treatment regime that can be individually tailored for each patient affected by neurodegenerative disorders. The polypeptides can also be used for diagnosing or treating Huntington's disease, Parkinson's disease, ADIS-related dementia or Tourette's syndrome. The present sequence represents a peptide that captured the survival-promoting activity of ADNF I, and is designated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New Activity Dependent Neurotrophic Factor I complex polypeptide, useful for reducing neuronal cell death, treating oxidative stress in a patient, or improving learning and/or memory in a subject with e.g. Alzheimer's
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10-APR-2002; 2002US-0371961P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 9 AA;
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                                                                                                   Homo sapiens
                                                                                                                               Human; glaucomatous optic neuropathy;
activity dependent neuroprotective factor;
                                                                                                                                                                         Human activity dependent neuroprotective factor (ADNP) peptide #1.
                                                                                                                                                                                                                                     ADA07952;
                                                                                                                                                                                                                                                                   ADA07952 standard; peptide; 9 AA
                                           04-SEP-2003
                                                                       US2003166544-A1
                                                                                                                                                                                                         20-NOV-2003
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DEPT HEALTH & HUMAN SERVICES.
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100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                      DB 6; L
                                                                                                                                  ADNP; ophthalmological
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06-JUN-2002; 2002US-00164432

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RESULT 9
ADC16629
ID C16629
ID ADC1
XX ADC1
XX ADC1
XX Huma
XX Heat
KW Inmu
KW Paras
KW Paras
KW Huma
XX Homc
KW Inmu
KW Paras
XX Homc
XX Hom
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  parasitic disease; inflammatory disease; autoimmune disease; graft rejection; cancer; allergy; Hsp60; antiinflammatory; immunosuppressive; antiparasitic; cytostatic; antiallergic; gene therapy;
                    Identifying a heat shock protein (Hsp) derived peptide for diagnosing or treating e.g. cancer by comparing the amount of cytokines produced by T cells exposed to antigen presenting cells not previously exposed to the test peptide.
                                                                                                                                                                                                                                                                    31-JAN-2002; 2002US-0352594P
                                                                                                                                                                                                                                                                                                                                                             07-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              heat shock protein; Hsp; antigen presenting cell; dendritic cell; immunomodulatory cytokine; cytokine; chemokine; surface antigen; parasitic disease; inflammatory disease; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC16629 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a method for preventing and treating glaucomatous optic neuropathy. The method comprises administering a composition comprising a peptide derived from activity dependent neuroprotective factor (ADNP). The method is useful for treating glaucomatous optic neuropathy. The present sequence represents a pept:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating glaucomatous optic neuropathy by administering a composition comprising a peptide derived from or related to Activity Dependent Neuroprotective Factor (ADNP).
                                                                                                                                                                                                                                                                                                              30-JAN-2003; 2003WO-IL000078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human heat shock protein 60 peptide p277 analogue Seq
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02-AUG-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 5; Page 2; 13pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-720933/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Clark AF, Shade DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CLAR/) CLARK A F. (SHAD/) SHADE D L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunosuppressive; a
numan; p277(442-450,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human ADNP.
                                                                                                                                    2003-663414/62.
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33.3%; Score 40; DB
Similarity 100.0%; Pred. No. 1.:
9; Conservative 0; Mismatches
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                                                                                                                                                                          Avron A,
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2001US-00921029.
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5. 1.8e+06;
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CC peptide for use in diagnosis or therapy. Exposure of antigen presenting CC cells, particularly dendritic cells, to peptides or peptide analogues CC derived from heat shock proteins subsequently activates T cells to produce immunomodulatory cytokines and will directly influence the CC cytokines, chemokines and surface antigens produced by the exposed CC antigen presenting cells. The present invention provides peptides and CC peptide analogues of heat shock proteins capable of directly interacting CC with dendritic cells. The peptides of the invention may be useful for the CC treatment of parasitic, inflammatory or autoimmune disease or graft CC rejection, cancer or allergy. The preferred peptides of the invention are CC peptide fragments (p277) or analogues (of p277), derived from heat shock protein Hep50. They may have antiallergic activities. The peptide CC sequences of the invention may also be useful for gene therapy. The CC (Hsp) 60 peptide fragment p277 analogue p277 (442-450, Ser6Ser11) of the CC invention
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Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                         immunosuppressive; nootropic; neuroprotective; antiinflammatory; vasotropic; muscular; CNS; thyromimetic; antithyroid; antirheumatic; antiarthritic; hepatotropic; virucide; dermatological; haemostatic; antidiabetic; antibacterial; neurotropic factor agonist; autoimmune disease; multiple sclerosis; mysathenia gravis; autoimmune disease; multiple sclerosis; mysathenia gravis; Guillain-Barre syndrome; systemic lupus erythematosus; Behcet's synd sjogren's syndrome; rheumatoid arthritis; Hashimoto's disease; synd sjogren's syndrome; rheumatoid arthritis; Hashimoto's disease; hypothyroiditis; primary biliary cirnhosis; mixed connective tissue disease; chronic active hepatitis; Graves' disease; hyperthyroiditis; scleroderma; chronic idiopathic thrombocytopenic purpura; diabetic neuropathy;
                                                                                                                      02-JAN-2003; 2003US-0437650P
                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADNF I active core site peptide SEQ ID NO:1.
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                                                                                                                                                                        30-DEC-2003; 2003WO-US041540
                                                                                                                                                                                                                                                                          WO2004060309-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           activity dependent neurotropic factor; ADNF; ADNF I; active core site;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel heat shock protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       41
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                                              UNIV RAM
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Offen D,
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                                              RAMOT AT TEL AVIV
EPT HEALTH & HUMAN
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Giladi E,
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1.8e+06;
  Brenneman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Behcet's syndrome;
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RESULT 11
ADS73608
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  active; core peptide; Activity Dependent Neurotrophic Factor; ADNF; ADNF II, anxiety; depression; hipophilic moiety; penetration; activity; panic disorder; occompulsive disorder; post-traumatic stress disorder; social phobia; social anxiety disorder; specific phobia; generalized anxiety disorder; Major depression; dysthymia; bipolar disorder; NAP-tubulin; binding site; anxiolytic drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    myasthenia gravis, Guillain-Barre syndrome, systemic lupus erythematosus, Behcet's syndrome, Sjogren's syndrome, rheumatoid arthritis, Hashimoto's disease/hypothyroiditis, primary biliary cirrhosis, mixed connective tissue disease, chronic active hepatitis, Graves' disease/hyperthyroiditis, scleroderma, chronic idiopathic thrombocytopenic purpura, diabetic neuropathy and septic shock. The present sequence represents an ADNF I active core site peptide from the
                                                                                                                                                                                                                             23-SEP-2004.
                                                                                                                                                                                                                                                                                                                                                                                 Homo
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                                                                         12-MAR-2003; 2003US-0454505P
                                                                                                                                               11-MAR-2004; 2004WO-IL000232
                                                                                                                                                                                                                                                                                                       WO2004080957-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                       neuroprotection.
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(UYRA-) UNIV RAMOT AT TEL AVIV LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       site.
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Pred. No. 1.8e+06;
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Example 4b;

Page 67; 121pp; English

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ARESULT 12
ARM64690
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ID ARM64690
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Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Activity dependent neurotrophic factor III: ADNF-III; ADNP; cell death; activity dependent neuroprotective protein; neurone; excito-toxicity; spinal cord; hippocampus; cerebral cortex; cholinergic; beta-amyloid; N-methyl-D-aspartate; Alzheimer's disease; human immunodeficiency virus
                                                  Activity dependent neurotrophic factor III polypeptide - therapeutically to prevent neuronal cell death associated infection, excito-toxicity or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV infection.
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                                                                                                                                                                                                                                                                                                                                                 07-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                       06-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-AUG-1998.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 SALLRSIPA 9
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Pred. No.
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Activity Dependent Neurotrophic Facto
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RESULT 13
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            epilepsy; AIDS dementia complex; neuropathic pain syndrome; AIS; amyotrophic lateral sclerosis; Parkinson's disease; Leber's disease; mitochondrial abnormality; Wernicke's encephalopathy; homocysteinuria; hyperprolinemia; sulphite oxide disease; Tourette's syndrome; notropic; Down's syndrome; drug addiction; developmental retardation; antilipemic; learning impairment; anticonvulsant; neuroprotective; anti-HIV.
                                                                                                     New nucleic acid encoding an activity (ADNF III) useful in the treatment of preventing neuronal cell death.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Activity Dependent Neurotrophic Factor III; ADNF; chromosome 20q13.2; Activity Dependent Neuroprotective Protein; ADNF; neuronal cell death; autosomal dominant nocturnal frontal-lobe epilepsy; ADNFLB; neurological deficiency; treatment; HIV; Human Immunodeficiency Virus; Alzheimer's disease; beta-amyloid peptide; Huntington's disease;
The present peptide sequence is used for affinity column chromatography, for purification of antibodies to Activity Dependent Neurotrophic Factor (ADNF), also called an Activity Dependent Neuroprotective Protein (ADNF)
                                                                                                                                                                             WPI; 2000-376491/32
                                                                                                                                                                                                                                                                                                       06-NOV-1998;
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                                                                    Example; Page 71; 136pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
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UNIV RAMOT APPLIED RES & IND DEV LTD.
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100.0%;
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                                                                                                                      dependent neurotrophic factor neurological deficiencies and
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RESULT 14
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            The present invention relates to the treatment of a condition associwith foetal alcohol syndrome (FAS), involving administering an action dependent neurotropic factor (ADNF). ADNFs of the present invention also be used to treat neurological deficiencies and prevent neuronal death. The present sequence is an ADNF peptide. (Updated on 14-MAY-2
                                                                                                                                                 exposed to alcadministering
                                                                                                                 Claim 6; Page 4; 65pp;
                                                                                                                                                                                  Treating
                                                                                                                                                                                                                      WPI; 2000-601940/57.
                                                                                                                                                                                                                                                      Brenneman DE,
                                                                                                                                                                                                                                                                                                                                       12-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neuronal
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                                                                                                                                                                                                                                                                                                         (UYRA-)
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                                                                                                                                                                                                                                                                                      ) UNIV RAMOT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N
                                                                                                                                                 g condition associated with fetal alcohol syndrome in a subject to alcohol in utero or reducing neuronal death, involves tering activity dependent neurotrophic factors I and/or III.
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Sequence 10

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RESULT 15
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                                                                                                              This invention relates to an activity dependent neurotrophic factor I (ADNF I) or ADNF III polypeptide. Sequences AAB72315 - AAB72321 represent CC ANNF II active core site peptides. Sequences AAB72322 - AAB72325 represent CC ADNF III active core site peptides. Sequences AAB72322 - AAB72326 represent CC composition containing either ADNF I or ADNF III and a pharmaceutical CC composition containing either ADNF I or ADNF III are useful for reducing CI neurons, cerebral cortical neurons and cholinergic neurons, hippocampal CI neurons, cerebral cortical neurons and cholinergic neurons, in a patient CI infected with a virus, e.g. human immunodeficiency virus (HIV). The CI neuronal cell death is associated with excito-toxicity induced by N-CI neuronal cell death is associated with excito-toxicity induced by N-CI neuronal cell death is associated with excito-toxicity induced by N-CI neuronal cell death is associated with excito-toxicity induced by Deta-amyloid CI neuronal cell death is associated with excito-toxicity induced by Deta-amyloid CI neuronal cell death is associated with excito-toxicity induced by Deta-amyloid CI neuronal cell death is associated with excito-toxicity induced by Deta-amyloid CI neuronal cell death is associated body weight, decreased by Cholinergic CI condition, such as decreased body weight, decreased brain weight, CI decreased level of vasoactive intestinal peptide (VIP) mRNA, and foetal CI death, associated with foetal alcohol syndrome
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Best Local Similarity 100.0%;
Matches 9; Conservative (
                      Query Match
Best Local S
      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel Activity Dependent Neurotrophic Factor I useful for treating oxidative stress, reducing neuronal cell death and treating a condition associated with fetal alcohol syndrome.
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                                                                                 Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 7; Page 56; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYRA-) UNIV RAMOT APPLIED RES & IND DEV LTD. (USSH ) US DEPT HEALTH & HUMAN SERVICES.
  Local Similarity les 9; Conserv
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      Conservative
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33.3%; bcc.
100.0%; Pr
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    Score 40; DB; Pred. No. 0.0
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VIP;
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